

Europäisches Patentamt European Patent Office Office européen des brevets



EP 1 201 765 A2

(12)

EUROPEAN PATENT APPLICATION

(43) Date of publication: 02.05.2002 Bulletin 2002/18

(21) Application number: 01124604.8

(22) Date of filing: 15.10.2001

(51) Int CI.7: **C12Q 1/48**, C07K 16/00, A61P 31/12, A61K 39/395, A61K 48/00, C12Q 1/68

(84) Designated Contracting States: AT BE CH CY DE DK ES FI FR GB GR IE IT LI LU MC NL PT SE TR Designated Extension States:

AL LT LV MK RO SI

(30) Priority: 16.10.2000 US 240750 P

(71) Applicant: Axxima Pharmaceuticals
Aktiengesellschaft
82152 Martinsried (DE)

(72) Inventors:

- Schubart, Daniel 79576 Weil am Rhein (DE)
- Habenberger, Peter 81373 Munich (DE)

 Stein-Gerlach, Matthias 81475 Munich (DE)

(11)

- Bevec, Dorian
 82110 Germering (DE)
- (74) Representative: Leidescher, Thomas et al Zimmermann & Partner, Postfach 33 09 20
 80069 München (DE)

Remarks:

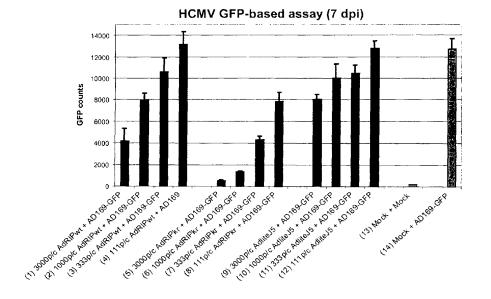
The sequence listing, which is published as annex to the application documents, was filed after the date of filing. The applicant has declared that it does not include matter which goes beyond the content of the application as filed.

(54) Cellular kinases involved in cytomegalovirus infection and their inhibition

(57) The role of certain cellular kinases active during Human Cytomegalovirus infection is disclosed. These cellular kinases are useful to detect HCMV infection,

and can be used to screen for cellular kinase inhibitors. Cellular kinases inhibitors, which effectively downregulate these key cellular components, serve as effective therapeutics against HCMV infection.

Fig. 1



Description

10

30

35

40

45

50

55

[0001] The present invention is in the fields of molecular biology and virology. The present invention is directed to novel methods for treating Cytomegalovirus using kinase inhibitors.

Background of the invention

[0002] Human Cytomegalovirus (HCMV) is a highly specific β-herpesvirus. Primary infection of healthy children and adults is usually asymptomatic, with a minority of cases developing a mononucleose-like syndrome. In contrast, congenital infection (U.S. 0.2%-2.2% per live birth; aprox. 40,000 per year) leads to several neurological defects in 10-15% of infected neonates. Immunocompromised patients represent another host group facing serious disease complications caused by HCMV infection or reactivation of a persistent infection. Up to 40% of the AIDS patients, for example, develop retinitis, pneumonitis, gastroenteritis or disseminated HCMV disease. In addition, allograft recipients (20,000 allograft transplantations per year in the U.S.) are often infected (or superinfected) by virus from the transplanted organ.

[0003] Clinical symptoms in the posttransplant period include prolonged fever, leukopenia, thrombocytopenia, atypical lymphocytosis, elevated hepatic transaminases and decreased graft survival. In bone marrow transplantations, HCMV infection is associated with high mortality rates (80-90% for untreated HCMV pneumonia).

[0004] Current approaches to develop therapeutics against Cytomegalovirus (CMV) have focused on antiviral agents per se; for example viral polymerase inhibitors. In fact, high mortality rates have been dramatically reduced by new antiviral agents. Current CMV therapeutics possess severe drawbacks, however. For example, Fomivirsen (Vitravene, formerly ISIS 2922) is typically administered by injection directly into the eye every 2 or 4 weeks. Ganciclovir is available for intravenous (Cytovene) or oral administration, and as an implant in the case of retinitis; unfortunately, toxic complications including leukopenia and thrombocytopenia frequently develop. Foscarnet (Foscavir; phosphonoformic acid), another antiviral agent, exhibits considerable renal toxicities and is only available in intravenous form (which is also true for Cidofovir (Vistide), another CMV therapeutic). In addition, CMV replication resumes soon after Ganciclovir and Foscarnet treatment is halted. Finally, Ganciclovir- and Foscarnet-resistant strains of CMV are emerging.

[0005] Although treatment of HCMV-induced disease has been improved with these inhibitors of the viral polymerase and preemptive or early antiviral therapy in transplant patients, there is a need in the art for a new class of HCMV therapeutics with better oral bioavailability and reduced toxic effects. This is especially true in the treatment of retinitis in AIDS patients, where CMV infection must be controlled for long periods of time.

[0006] Recent research has revealed how cells communicate with each other to coordinate the growth and maintenance of the multitude of tissues within the human body. A key element of this communication network is the transmission of a signal from the exterior of a cell to its nucleus, which results in the activation or suppression of specific genes. This process is called signal transduction.

[0007] An integral part of signal transduction is the interaction of cytokines, their receptors, and intracellular signal transduction molecules. Cytokines serve as messengers that bind to receptors on the surface of a target cell. As a result of the binding, the receptors activate a cascade of downstream signaling molecules, thereby transmitting the message from the exterior of the cell to its nucleus. Signal transduction to the nucleus modulates specific gene expression (i.e., transcription and translation), which results in either the upregulation or downregulation of specific proteins that carry out a particular biological function.

[0008] Viral infection disrupts normal signal transduction, which leads to cellular malfunctioning resulting in a disease state. Specifically, interference of HCMV with relevant human primary cells is necessary for the virus to create an environment that allows it to grow and replicate, and in turn cause disease in the infected individual. Current research efforts have failed to elucidate all the specific intracellular signal pathways affected by HCMV infection, however. Discovery of the signal transduction pathways and specific intracellular signal transduction molecules affected by CMV infection would represent a tremendous advance in the understanding of the induction and progression of CMV infection processes and provide new avenues for the development of a novel class of effective therapeutics for the treatment of CMV.

[0009] Thus, object of the present invention is to provide methods for detecting, preventing and/or treating Cytomegalovirus infection and/or associated diseases, methods for the identification of compounds useful for preventing and/or treating Cytomegalovirus infection and/or associated diseases and for regulating the production of Cytomegaloviruses.

[0010] The object of the present invention is solved by the teaching of the independent claims. Further advantageous features, aspects and details of the invention are evident from the dependent claims, the description, and the examples of the present application.

Description of the invention

10

20

25

30

35

40

45

50

[0011] The present invention is based upon the finding of a group of cellular kinases that are specifically upregulated as a result of CMV infection. The antiviral therapeutic research approach described herein, focuses on discovering the cellular signal transduction pathways involved in viral infection. Identification of the cellular signal transduction molecules, key to viral infection provides for, among other things, novel diagnostic methods, especially assays, and compositions useful therefor, novel targets for antiviral therapeutics, a novel class of antiviral therapeutics, and new screening assays and materials to discover new antiviral agents.

[0012] This approach led to the development of a novel microarray platform technology, wherein a microarray of more than 1100 signal transduction cDNAs was developed. This unique microarray technology was used to identify RNA expression patterns (e.g., upregulation or downregulation) unique to CMV infected host cells. Differential display techniques were used to pinpoint those signal transduction molecules useful as targets for drug intervention. Effective manipulation of these virally-controlled intracellular signal transduction pathways can alter (slow or stop altogether) the course of viral growth.

[0013] It is now revealed for the first time that the cellular protein kinases RICK (also known as CARDIAK; RIP2), RIP, NIK (also known as HGK; MAP4K4), MKK3 (also known as MEK3), and SRPK-2 are specifically and uniquely upregulated in a cell as a result of CMV infection. These cellular kinases therefore identify novel diagnostic and therapeutic targets for CMV infection.

[0014] Surprisingly, it was found that the following human cellular targets are significantly upregulated compared with uninfected human foreskin fibroblasts cells:

target	upregulation
RICK	3.6 fold
RIP	2.6 fold
NIK	4.0 fold
МККЗ	2.5 fold
SRPK-2	2.2 fold

[0015] Based upon the research work reported herein, one aspect of the present invention is directed to a method, preferably a screening assay, for identifying compounds useful for treating and/or preventing Cytomegalovirus infection and/or diseases associated therewith. Specifically, this assay involves contacting a test compound with one or more cellular kinases selected from the group consisting of RICK, RIP, NIK, MKK3, and SRPK-2, and detecting a change, normally a decrease, in activity of said cellular kinase. This method was used for the identification of the RICK and RIP inhibitors shown below in Table 1 and Table 2.

[0016] Another aspect of the invention is directed to a diagnostic method for detecting Cytomegalovirus infection and/or associated diseases in an individual or in cells and/or in cell lysates. This assay involves providing a sample from the individual or providing a sample from said cells, respectively, and detecting activity of one or more cellular kinases selected from the group consisting of RICK, RIP, NIK, MKK3, and SRPK-2. The term "individual" preferably refers to mammals, especially humans or ruminants.

[0017] Also described in the present invention are monoclonal or polyclonal antibodies which bind to a cellular kinase selected from the group consisting of RICK, RIP, NIK, MKK3, and SRPK-2.

[0018] A further aspect of the present invention relates to a method for preventing and/or treating Cytomegalovirus infection and/or associated diseases in an individual by administering a pharmaceutically effective amount of an inhibitor to said individual, wherein said inhibitor inhibits at least partially the activity of one or more cellular kinases selected from the group consisting of RICK, RIP, NIK, MKK3, and SRPK-2, or wherein said inhibitor inhibits at least partially the production of one or more cellular kinases selected from the group consisting of RICK, RIP, NIK, MKK3, and SRPK-2. [0019] As used herein, the term "inhibitor" refers to any compound capable of downregulating, decreasing, reducing, suppressing or inactivating the amount and/or activity of at least one human cellular protein kinase selected from the group consisting of RICK, RIP, NIK, MKK3, and SRPK-2. Generally, said inhibitors, including suicide inhibitors, may be proteins, oligo- and polypeptides, nucleic acids, genes, small chemical molecules, or other chemical moieties. Suitable inhibitors are monoclonal or polyclonal antibodies which bind to at least one cellular kinase selected from the group consisting of RICK, RIP, NIK, MKK3, and SRPK-2.

[0020] Based on the surprising results reported herein, one aspect of the present invention is directed to a method for regulating the production of Cytomegalovirus in an individual by administering an individual a pharmaceutically effective amount of an inhibitor wherein said inhibitor inhibits at least partially the activity of one or more cellular kinases selected from the group consisting of RICK, RIP, NIK, MKK3, and SRPK-2, or wherein said inhibitor at least partially

inhibits the production of one or more cellular kinases selected from the group consisting of RICK, RIP, NIK, MKK3, and SRPK-2.

[0021] A similar aspect relates to a method for regulating the production of Cytomegalovirus in cells by administering the cells a pharmaceutically effective amount of an inhibitor wherein said inhibitor inhibits at least partially the activity of one or more cellular kinases selected from the group consisting of RICK, RIP, NIK, MKK3, and SRPK-2, or wherein said inhibitor at least partially inhibits the production of one or more cellular kinases selected from the group consisting of RICK, RIP, NIK, MKK3, and SRPK-2 in the cells.

[0022] Yet another aspect of the invention is directed to a method for regulating the expression of at least one cellular kinase selected from the group consisting of RICK, RIP, NIK, MKK3, and SRPK-2 in an individual comprising the step of administering the individual a pharmaceutically effective amount of an inhibitor wherein said inhibitor inhibits at least partially the transcription of DNA or the translation of RNA encoding one of said cellular kinases.

10

30

35

40

50

[0023] A further aspect relates to a method for regulating the expression of at least one cellular kinase selected from the group consisting of RICK, RIP, NIK, MKK3, and SRPK-2 in the cells comprising the step of administering the cells a pharmaceutically effective amount of an inhibitor wherein said inhibitor inhibits at least partially the transcription of DNA or the translation of RNA encoding one of said cellular kinases.

[0024] As used herein, the term "regulating expression and/or activity" generally refers to any process that functions to control or modulate the quantity or activity (functionality) of a cellular component. Static regulation maintains expression and/or activity at some given level. Upregulation refers to a relative increase in expression and/or activity. Accordingly downregulation refers to a relative decrease in expression and/or activity. In the present invention, regulation is preferably the downregulation of a cellular component. Downregulation is synonymous with inhibition of a given cellular component's activity.

[0025] Beside inhibitors also activators may be useful for treating Cytomegalovirus infection by increasing the activity of at least one of the cellular protein kinases RICK, RIP, NIK, MKK3, and SRPK-2. Thus, a method for preventing and/or treating Cytomegalovirus infection and/or associated diseases in an individual is disclosed. Said method comprises administering a pharmaceutically effective amount of an activator to an individual, wherein said activator activates at least partially the activity of one or more cellular kinases selected from the group consisting of RICK, RIP, NIK, MKK3, and SRPK-2, or wherein said activator activates or stimulates at least partially the production of one or more cellular kinases selected from the group consisting of RICK, RIP, NIK, MKK3, and SRPK-2.

[0026] Furthermore, a method for regulating the production of Cytomegalovirus either in cells or in an individual is described. Said methods comprise administering an individual or to cells a pharmaceutically effective amount of an activator wherein said activator activates at least partially the activity of one or more cellular kinases selected from the group consisting of RICK, RIP, NIK, MKK3, and SRPK-2, or wherein said activator at least partially activates or stimulates the production of one or more cellular kinases selected from the group consisting of RICK, RIP, NIK, MKK3, and SRPK-2.

[0027] As used herein, the term "activator" refers to any chemical compound which is able to upregulate, increase, activate, or stimulate the activity of at least one human cellular protein kinase selected from the group consisting of RICK, RIP, NIK, MKK3, and SRPK-2 or which is able to upregulate, increase, activate, or stimulate the expression of at least one of said cellular kinases. Activators comprise proteins, oligo- and polypeptides, nucleic acids, genes, and preferably small chemical molecules, or other chemical moieties.

[0028] Still another aspect of the present invention is directed to either a method for regulating the expression of at least one cellular kinase selected from the group consisting of RICK, RIP, NIK, MKK3, and SRPK-2 in an individual or for regulating the expression of at least one of said kinases in cells. These methods comprise the step of administering the individual or the cells a pharmaceutically effective amount of an activator wherein said activator activates at least partially the transcription of DNA or the translation of RNA encoding one of said cellular kinases.

[0029] Furthermore, oligonucleotides are disclosed which bind to the DNA or RNA encoding a cellular kinase selected from the group consisting of RICK, RIP, NIK, MKK3, and SRPK-2. Said oligonucleotides can be used as suitable inhibitors within the aforementioned methods.

[0030] Some methods of the present invention identify compounds useful for prophylaxis and/or treatment of Cytomegalovirus infection and/or associated disease by screening a test compound, or a library of test compounds, for its ability to inhibit at least one of the above-mentioned human cellular protein kinases identified herein as characteristically upregulated during HCMV replication. Using this method the compounds A to E have been identified as RICK inhibitors and the compounds F to H have been identified as RIP inhibitors. Thus, the use of these compounds as inhibitors of RICK or RIP is disclosed. Furthermore, these compounds can be used for manufacturing a pharmaceutical composition for prophylaxis and/or treatment of Cytomegalovirus infection and/or diseases associated with Cytomegalovirus infection.

[0031] A variety of assay protocols and detection techniques are well known in the art and easily adapted for this purpose by a skilled practitioner. Such methods include, but are not limited to, high throughput assays (e.g., microarray technology, phage display technology), and *in vitro* and *in vivo* cellular and tissue assays.

[0032] Thus, some embodiments of the present invention may comprise a solid support useful for detecting Cytomegalovirus infection in a cell or an individual. Preferably the solid support comprises immobilized oligonucleotides, wherein the oligonucleotides are capable of detecting activity of one or more cellular kinases selected from the group consisting of: RICK, RIP, NIK, MKK3, and SRPK-2.

[0033] Another aspect of the invention includes a solid support useful for screening compounds useful for treating Cytomegalovirus. Preferred embodiments include a solid support comprising one or more immobilized oligonucleotides, wherein the oligonucleotide(s) encode one or more cellular kinases selected from the group consisting of RICK, RIP, NIK, MKK3, and SRPK-2. In another preferred embodiment, the solid support comprises one or more immobilized cellular kinases selected from the group consisting of: RICK, RIP, NIK, MKK3, and SRPK-2.

[0034] Accordingly, another aspect of the present invention is directed to a novel therapeutic composition useful to treat an individual afflicted with Cytomegalovirus comprising one or more inhibitors capable of inhibiting activity of one or more of the cellular kinases selected from the group consisting of RICK, RIP, NIK, MKK3, and SRPK-2. In addition thereto, a novel pharmaceutical composition could comprise at least one inhibitor capable of regulating the production of HCMV by inhibiting the expression of at least one cellular kinase selected from the group consisting of RICK, RIP, NIK, MKK3, and SRPK-2.

[0035] Another group of suitable therapeutic compositions useful for prophylaxis and/or treatment of CMV comprises at least one activator which is able to increase the activity of one or more cellular kinases selected from the group consisting of RICK, RIP, NIK, MKK3, and SRPK-2 or which is capable of increasing the expression of at least one cellular kinase selected from the group consisting of RICK, RIP, NIK, MKK3, and SRPK-2.

[0036] Said pharmaceutical compositions may further comprise pharmaceutically acceptable carriers, excipient, diluents, fillers, binders, disintegrants, lubricants, glidents, coloring agents, flavoring agents, opaquing agents, and/or plasticizers.

Detailed description of the invention

[0037] Utilizing microarray technology, a unique microarray of more than 1100 signal transduction cDNAs was developed. This array was used to compare signal transduction mRNA expression patterns (e.g., upregulation or down-regulation) in primary human cells before and after infection with HCMV at various timepoints of infection. Interference of the HCMV with the cellular signaling events is reflected in differential gene expression when compared to the uninfected cellular signaling. Results from this novel signal transduction microarray analysis revealed significant upregulation of cellular protein kinases RICK, RIP, NIK, MKK3, and SRPK-2, as unique to CMV infected host cells. These findings were confirmed utilizing conventional Northern and Western blot analyses.

[0038] Disclosed herein is the first report describing the role of cellular kinases; RICK, RIP, NIK, MKK3, and SRPK-2 in the signal transduction of CMV viral infection process. As a result of these discoveries, a novel class of compounds, i.e., RICK, RIP, NIK, MKK3, and SRPK-2 inhibitors, are identified as useful for altering the course of CMV infection.

[0039] To perform initial tests for compounds that inhibit RICK activity in a cellular assay, RICK was transiently overexpressed in HEK-293 cells, immunoprecipitated and incubated with different concentrations of test compounds before in-vitro kinase assays were performed (Example 10). According to the method for identifying compounds useful for inhibiting the cellular kinase RICK and therefore useful for treating and/or preventing Cytomegalovirus infection and/ or diseases associated with Cytomegalovirus infection, a test compound is contacted with the cellular kinase RICK according to the RICK assay protocoll disclosed in example 10. The test compound dissolved in DMSO is added to the RICK assay solution at concentrations between 100 nM and 50 μM. Thereafter, radioactively labeled ATP is added and kinase activity of RICK is determined by detecting the autophosphorylation of RICK via radioactivity measurement. The five compounds listed in the following Table 1 were identified using said method. These compounds showed inhibition of RICK kinase activity with an IC $_{50}$ between about 500 nM and 1 μ M and an inhibition of HCMV with an IC $_{50}$ between about 1 and 8 μ M, respectively. The IC₅₀ values of HCMV inhibition were obtained by the use of at least one assay protocol selected from a) virus replication assay, b) plaque assay, c) GFP (Green Fluorescent Protein) infection assay, and d) indirect immunofluorescence analysis as disclosed in example 12. Thus, the five compounds A to E mentioned below and/or pharmaceutically acceptable salts thereof can be used as inhibitors of the cellular protein kinase RICK and as pharmaceutically active compounds for the treatment and/or prophylaxis of HCMV infection. Furthermore, these compounds are suitable for the manufacture of a pharmaceutical composition for prophylaxis and/or treatment of Cytomegalovirus infection and/or diseases associated therewith.

55

10

25

30

35

40

45

50

Table 1: Inhibitors of RICK and HCMV

5	compound	structure	IC ₅₀ RICK	IC ₅₀ HCMV
10	A \	CI N N N CH ₃	1 μΜ	6.8 µM
20	B	N CH ₃	500 nM	1.4 μΜ
30	H ₃ C N N	CH ₃	1 μΜ	6.2 μM
35				

5	H₃C H₃C	-O N N N Br	500 nM	7.6 µM
15	E	H ₃ CO N N N N N N N N N N N N N N N N N N N	500 nM	5.7 μ M

[0040] The compounds A to E have the following names:

Compound A: 6-(2,6-Dichlorophenyl)-8-methyl-2-(3-morpholin-4-yl-propylamino)-8*H*-pyrido[2,3-*a*]pyrimidin-7-one;

Compound B: 8-methyl-6-phenyl-2-(pyridin-4-yl-amino)-8*H*-pyrido[2,3-*d*]pyrimidin-7-one;

Compound C: 6-(2,6-Dichlorophenyl)-8-methyl-2-[3-(4-methylpiperazin-1-yl)-propylamino]-8H-pyrido[2,3-d]pyrimi-

din-7-one;

25

30

50

Compound D: (3-Bromophenyl)-(6,7-diethoxyquinazolin-4-yl)-amine; Compound E: (3-Bromophenyl)-(6,7-dimethoxyquinazolin-4-yl)-amine.

[0041] From the results observed with the compounds shown in Table 1 it is proved that RICK is an important target for the treatment of HCMV and diseases associated with HCMV infection. Inhibitors of the human cellular protein kinase RICK may serve as new pharmaceutical substances for prophylaxis and/or treatment of Cytomegalovirus infection and/or diseases associated with CMV infection.

[0042] In addition to the chemical validation of RICK described above, a genetic validation of RICK in HCMV infection was performed. Wildtype and mutated RICK was expressed in HFF cells with a modified Adenovirus as vehicle (Example 9). The expression of both, wildtype and mutated RICK, caused a dramatic reduction in HCMV replication (cf. Fig. 2). Also these data confirm RICK as a valuable therapeutic target in HCMV treatment. As known in the art and as used herein, "RICK" refers to a protein kinase also known as "CARDIAK" and as "RIP2" which is a RIP-like kinase. RICK is essentially characterized as comprising an N-terminal serine-threonine kinase catalytic domain and a C-terminal region containing a caspase-recruitment domain (referred to as "CARD").

[0043] To perform initial tests for compounds that inhibit RIP activity in a cellular assay, RIP was transiently overexpressed in HEK-293 cells, immunoprecipitated and incubated with different concentrations of test compounds before in-vitro kinase assays were performed (Example 11). In order to identify compounds suitable for inhibiting the cellular kinase RIP and, thus, suitable for treating and/or preventing Cytomegalovirus infection and/or diseases associated with Cytomegalovirus infection the inventive method according to claim 1 was used. A test compound was contacted with the cellular kinase RIP according to the RIP assay protocoll disclosed in example 11. The test compound dissolved in DMSO is added to the RIP assay solution at concentrations between 100 nM and 50 μ M. Radioactively labeled ATP was used as co-substrate of RIP and auto-phosphorylation was detected via measurement of incorporation of radioactivity into the RIP protein. Thereafter, phosphorylation rates with and without test compounds were compared. The three compounds listed in the following Table 2 showed inhibition of RIP kinase activity with an IC $_{50}$ between about 5 μ M and 10 μ M and an inhibition of HCMV with an IC $_{50}$ between about 12 μ M and 15 μ M, respectively. The IC $_{50}$ values of HCMV inhibition were obtained by the use of at least one assay protocol selected from a) virus replication assay, b) plaque assay, c) GFP infection assay, and d) indirect immunofluorescence analysis as disclosed in example 12.

[0044] Thus, the three compounds F to H mentioned below and/or pharmaceutically acceptable salts thereof can be used as inhibitors of the cellular protein kinase RIP and as pharmaceutically active compounds for the treatment and/or prophylaxis of HCMV infection. Furthermore, these compounds are suitable for the manufacture of a pharmaceutical composition for prophylaxis and/or treatment of Cytomegalovirus infection and/or diseases associated therewith.

Table 2: Inhibitors of RIP and HCMV

5

10

15

20

25

30

35

40

45

50

55

compound	structure	IC ₅₀ RIP	IC ₅₀ HCMV
но	OH OH	5 µM	15 µM
Ga	N O N	10 µM	15 µM
Н	N O N O	5 µM	12 μΜ

[0045] The compounds F to H have the following names:

Compound F: 2-(3,4-Dihydroxyphenyl)-3,5,7-trihydroxychromen-4-one; Compound G: 5-Cloro-3-(1*H*-pyrrol-2-vlmethylene)-1,3-dihydroindol-2-one;

Compound H: 4-Quinolin-4-ylmethylene-4*H*-isoquinoline-1,3-dione.

[0046] From the results observed with the compounds shown in Table 2 it is proved that RIP is an important target for the treatment of HCMV and diseases associated with HCMV infection. Inhibitors of the human cellular protein kinase RIP may serve as new pharmaceutical substances for prophylaxis and/or treatment of Cytomegalovirus infection and/or diseases associated with CMV infection.

[0047] In addition to the chemical validation of RIP described above, a genetic validation of RIP in HCMV infection was performed. Wildtype and mutated RIP was expressed in HFF cells with a modified Adenovirus as vehicle (Example 9). The expression of mutated RICK, but not wildtype RIP, caused a dramatic reduction in HCMV replication (cf. Fig. 1). These data also confirm RIP as therapeutic target in HCMV treatment.

[0048] As known in the art and as used herein, "NIK" (Nck-Interacting Kinase; also known as "HGK" or "MAP4K4") refers to an NF-kappaB inducing serine/threonine kinase that interacts with the SH3 domains of Nck (an adaptor protein composed of one SH2 and three SH3 domains, known as a common target for a variety of cell surface receptors). NIK is most homologous to the Sterile 20 (Ste20) family of protein kinases, particularly GCK and MSST1 in that they bind

neither Cdc42 nor Rac and contain an N-terminal kinase domain with a putative C-terminal regulatory domain. NIK is reported to promote neurite process formation and mediated anti-apoptotic signaling. NIK expression leads to IKK activation and induced nuclear translocation of NF-kappaB. NIK activates MEK1 phosphorylation and induces the Erk1/Erk2 MAPK pathway. NIK has been shown to be a MEK1-dependent activator of the MAPK pathway (Foehr et al., 2000. J. Biol. Chem. 275, 34021-34024). Overexpression of NIK has been reported to specifically activate the stress-activated protein kinase (SAPK) pathway; possibly upstream of MEKK1, a dominant-negative MEK kinase 1 capable of blocking NIK activation of SAPK (Su et al., 1997. EMBO 16(6):1279-90).

[0049] As known in the art and as used herein, "MKK3" (MAP kinase kinase 3; also known as "MEK3") refers to a protein kinase known to function in TNF-induced cytokine expression, and specifically phosphorylate and activate p38 MAP kinase (Blank et al., 1996. <u>J. Biol. Chem.</u> 271:5361-5368; Raingeaud et al., 1996. <u>Mol. Cell. Biol.</u> 16(3):1247-55). MKK3 gene disruption has been shown to cause a selective defect in the response of fibroblasts to the proinflammatory cytokine tumor necrosis factor, including reduced p38 MAP kinase activation and cytokine expression; suggesting that the MKK3 protein kinase is a critical component of a tumor necrosis factor-stimulated signaling pathway that causes increased expression of inflammatory cytokines (Wysk et al., 1999. PNAS USA 96(7):3763-8).

10

30

35

40

50

[0050] As known in the art and as used herein, "SRPK-2" (SR-protein-specific kinase 2) refers to a kinase known to phosphorylate SF2/ASF and believed to regulate the disassembly of the SR family of splicing factors in a tissue-specific manner (e.g., in testis, lung, and brain; Kuroyanagi et al., 1998. <u>Biochem. Biophys. Res. Commun.</u> 242(2):357-64). SRPK-2 is believed to function in spliceosome assembly and in mediating the trafficking of splicing factors (Wang et al., 1998. J. Cell. Biol. 140(4):737-50; Wang et al., 1999. Genomics 57(2):310-5).

[0051] In one embodiment, the present invention is directed to a method for treating CMV infection by administering a pharmaceutically effective amount of an inhibitor of one or more of the cellular kinases; RICK, RIP, NIK, MKK3, and/or SRPK-2.

[0052] As used herein, a cellular kinase "inhibitor" refers to any compound capable of downregulating, decreasing, suppressing or otherwise regulating the amount and/or activity of a cellular kinase. Inhibition of these cellular kinases can be achieved by any of a variety of mechanisms known in the art, including, but not limited to binding directly to the cellular kinase polypeptide (e.g., a RICK-inhibitor compound binding complex, or substrate mimetic), denaturing or otherwise inactivating the cellular kinase, or inhibiting the expression of the gene (e.g., transcription to mRNA, translation to a nascent polypeptide, and/or final polypeptide modifications to a mature protein), which encodes the cellular kinase. Generally, cellular kinase inhibitors may be proteins, polypeptides, nucleic acids, small molecules, or other chemical moieties.

[0053] Yet another aspect of the present invention is directed to pharmaceutical compositions useful for the prophylaxis and/or treatment of an individual afflicted with Cytomegalovirus infection and/or associated diseases. Said pharmaceutical composition comprises at least one pharmaceutically active compound capable of regulating at least partially the activity or the expression of one human cellular protein kinase selected from the group comprising RICK, RIP, NIK, MKK3, and SRPK-2 and/or capable of regulating the replication of CMV.

[0054] As used herein the term "regulating" refers either to the ability of an inhibitor to downregulate, decrease, reduce, suppress, inactivate, or inhibit at least partially the activity of an enzyme, or the expression of an enzyme and the virus replication or to the ability of an activator to upregulate, increase, stimulate, or activate at least partially the activity of an enzyme or the expression of an enzyme.

[0055] Suitable examples for inhibitors which are the pharmaceutically active components within the therapeutic compositions are the compounds A to H mentioned in Table 1 and 2. The compounds 6-(2,6-Dichlorophenyl)-8-methyl-2-(3-morpholin-4-yl-propylamino)-8H-pyrido[2,3-d]pyrimidin-7-one; 8-methyl-6-phenyl-2-(pyridin-4-yl-amino)-8H-pyrido[2,3-d]pyrimidin-7-one; 6-(2,6-Dichlorophenyl)-8-methyl-2-[3-(4-methylpiperazin-1-yl)-propylamino]-8H-pyrido [2,3-d]pyrimidin-7-one; (3-Bromophenyl)-(6,7-dimethoxyquinazolin-4-yl)-amine; (3-Bromophenyl)-(6,7-diethoxyquinazolin-4-yl)-amine; 2-(3,4-Dihydroxyphenyl)-3,5,7-trihydroxychromen-4-one; 5-Cloro-3-(1H-pyrrol-2-ylmethylene)-1,3-dihydroindol-2-one; 4-Quinolin-4-ylmethylene-4H-isoquinoline-1,3-dione and/or pharmaceutically acceptable salts of these compounds are useful for prophylaxis and/or treatment of Cytomegalovirus infection and/or diseases associated with Cytomegalovirus infection.

[0056] CMV therapeutics may be administered to cells from an individual *in vitro*, or may involve *in vivo* administration to the individual. Routes of administration of pharmaceutical preparations to an individual may include inhalation, oral and parenteral, including dermal, intradermal, intragastral, intracutan, intravasal, intravenous, intramuscular, intraperitoneal, intranasal, intravaginal, intrabuccal, percutan, rectal, subcutaneous, sublingual, topical or transdermal application, but are not limited the these ways of administration. For instance, the preferred preparations are in administratable form which is suitable for oral application. These administratable forms, for example, include pills, tablets, film tablets, coated tablets, capsules, powders and deposits. Administration to an individual may be in a single dose or in repeated administrations, and may be in any of a variety of physiologically acceptable salt forms, and/or with an acceptable pharmaceutical carrier, binder, lubricant, excipient, diluents and/or adjuvant. Pharmaceutically acceptable salt forms and standard pharmaceutical formulation techniques are well known to persons skilled in the art (see, for

example, Remington's Pharmaceutical Sciences, Mack Publishing Co.).

10

30

35

40

45

50

[0057] As used herein, a "pharmaceutically effective amount" of a cellular kinase inhibitor is an amount effective to achieve the desired physiological result, either in cells treated in vitro or in a subject treated in vivo. Specifically, a pharmaceutically effective amount is an amount sufficient to inhibit, for some period of time, one or more of the clinically defined pathological processes associated with the viral infection. The effective amount may vary depending on the specific kinase inhibitor selected, and is also dependent on a variety of factors and conditions related to the subject to be treated and the severity of the infection. For example, if the inhibitor is to be administered in vivo, factors such as the age, weight and health of the patient as well as dose response curves and toxicity data obtained in preclinical animal work would be among those considered. If the inhibitor is to be contacted with the cells in vitro, one would also design a variety of pre-clinical in vitro studies to assess such parameters as uptake, half-life, dose, toxicity, etc. The determination of a pharmaceutically effective amount for a given agent is well within the ability of those skilled in the art. [0058] As a result of the discovery of the upregulation of certain cellular kinases as part of the infection process of CMV, a novel diagnostic assay useful for the detecting CMV infection of an individual (or cell) is identified. The diagnostic assay of the present invention involves providing a sample from an individual or providing cells and/or cell lysates, and detecting activity in the sample of one or more cellular kinases selected from the group consisting of RICK, RIP, NIK, MKK3, and SRPK-2. In one embodiment, deviations in the expression levels of one or more of the identified cellular kinases in a test sample compared to a known normal expression levels (e.g., determined from a sample from a healthy individual) will be diagnostic of CMV.

[0059] It is apparent to a practitioner in the art that a sample useful for detecting CMV infection, whether of a subject individual or an isolated cell, refers to any cellular extract (including whole cells) from a tissue or body fluid (in the case of an individual) or cellular lysate (in the case of an isolated cell), which contains cellular components representative of cellular activity of one or more of the above-mentioned cellular kinases.

[0060] It is also apparent to a person of ordinary skill in the art that detection includes any method known in the art useful to indicate the presence, absence, or amount of a detection target. Such methods may include, but are not limited to, any molecular or cellular techniques, used singularly or in combination, including, but not limited to: hybridization and/or binding techniques, including blotting techniques and immunoassays; labeling techniques (chemiluminescent, colorimetric, fluorescent, radioisotopic); spectroscopic techniques; separations technology, including precipitations, electrophoresis, chromatography, centrifugation, ultrafiltration, cell sorting; and enzymatic manipulations (e. q., digestion).

[0061] Because the present disclosure teaches for the first time the upregulation of a group of cellular kinases specifically involved in the viral infection of CMV, the present invention is also directed to an assay useful for detecting novel compounds useful for treating CMV infection.

[0062] Assays of the present invention identify compounds useful for treating CMV operate by screening a test compound, or library of test compounds, for its ability to inhibit any one or more of the group of cellular kinases identified herein as characteristically upregulated during CMV growth and replication inside a cell. A variety of assay protocols and detection techniques are well known in the art and easily adapted for this purpose by a skilled practitioner. Such assays include, but are not limited to, high throughput assays (e.g., microarray technology, phage display technology), and *in vitro* and *in vitro* cellular and tissue assays.

[0063] In a related aspect, it is also an object of the present invention, in view of the discovery of cellular kinases specifically involved in CMV growth in a cell, to provide an assay component specially useful for detecting CMV in an individual (or a cell). Preferably the assay component comprises oligonucleotides capable of detecting activity of one or more of the cellular kinases RICK, RIP, NIK, MKK3, and SRPK-2 in a sample (e.g., by hybridization to mRNA from the sample), immobilized on a solid support. Most preferably the solid support would contain oligonucleotides of sufficient quality and quantity to detect all of the above-mentioned cellular kinases (e.g., a nucleic acid microarray).

[0064] Similarly, it is part of the object of the invention to provide an assay component specially useful for screening compounds useful for treating CMV. One preferred assay component comprises oligonucleotides that encode one or more of the cellular kinases RICK, RIP, NIK, MKK3, and SRPK-2, immobilized on a solid support. In another embodiment, the assay component comprises peptide fragments of one or more of the above-identified cellular kinases immobilized on a solid support. Once again the most preferred solid support embodiment would contain polymers of sufficient quality and quantity to detect all of the above-mentioned cellular kinases (e.g., a nucleic acid or a peptide microarray). A variety of assay supports and construction of the same are well known in the art and easily adapted for this purpose by a skilled practitioner (see, for example: Marshall, 1999. "Do-it-yourself gene watching" Science 286: 444-447 (including insets); and Service, 2000. "Protein arrays step out of DNA's shadow" Science 289:1673).

[0065] It is preferred that mRNA is assayed as an indication of expression. Methods for assaying for mRNA include, but are not limited to, Northern blots, slot blots, dot blots, and hybridization to an ordered array of oligonucleotides. Nucleic acid probes useful for assay of a sample are preferably of sufficient length to specifically hybridize only to appropriate, complementary transcripts. Typically the oligonucleotide probes will be at least 10, 12, 14, 16, 18, 20 or 25 nucleotides in length. In some cases longer probes of at least 30, 40, or 50 nucleotides will be desirable.

[0066] The cDNA oligonucleotides immobilized on said membrane filter which are used for detecting the up- or down-regulation of the above-mentioned human cellular protein kinases by hybridization to the radioactively labeled cDNA probes have the nucleotide sequences listed in table 3.

Table 3:

5

10

15

30

35

40

45

50

Nucleotide sequences of cDNA-arrays											
Human cellular kinase Sequence of immobilized DNA on arrays (in relation to the respective Acc I											
RICK	914 bp - 2501 bp (AF027706)										
RIP	1421 bp - 2617 bp (U50062)										
NIK	231 bp - 3077 bp (Y10256)										
МККЗ	341 bp - 2030 bp (NM_002756)										
SRPK-2	1238 bp - 2790 bp (U88666)										

[0067] The nucleoside sequences of the genes coding for the human cellular protein kinases RICK, RIP, NIK, MKK3, and SRPK-2 listed in Table 3 together with the amino acid sequences of said enzymes can be obtained from NCBI (National Library of Medicine: PubMed; Web address: www.ncbi.nlm.nih.gov/entrez). Sequence protocols of the five cellular protein kinases are attached to this application as a part of the description.

[0068] The polypeptide product of gene expression may be assayed to determine the amount of expression as well. Methods for assaying for a protein include, but are not limited to, Western blot, immunprecipitation, radioimmunoassay and peptide immobilization in an ordered array. It is understood, however, that any method for specifically and quantitatively measuring a specific protein or mRNA product can be used.

[0069] A variety of supports upon which nucleic acids or peptides can be immobilized are known in the art, for example filters, or polyvinyl chloride dishes. Any solid surface to which oligonucleotides or peptides can be bound, either directly or indirectly, either covalently or non-covalently, can be used. A preferred solid support is a microarray membrane filter or a "biochip". These contain particular polymer probes in predetermined locations on the array. Each predetermined location may contain more than one molecule of the probe, but each molecule within the predetermined location has an identical sequence.

[0070] The present invention incorporates by reference in their entirety techniques well known in the field of molecular biology. These techniques include, but are not limited to, techniques described in the following publications:

Ausubel, F.M. et al. eds., Short Protocols In Molecular Biology (4th Ed. 1999) John Wiley & Sons, NY. (ISBN 0-471-32938-X).

Old, R.W. & S.B. Primrose, Principles of Gene Manipulation: An Introduction To Genetic Engineering (3d Ed. 1985) Blackwell Scientific Publications, Boston. Studies in Microbiology; V.2:409 pp. (ISBN 0-632-01318-4).

Miller, J.H. & M.P. Calos eds., <u>Gene Transfer Vectors For Mammalian Cells</u> (1987) Cold Spring Harbor Laboratory Press, NY. 169 pp. (ISBN 0-87969-198-0).

Mayer, R.J. & J.H. Walker eds., <u>Immunochemical Methods In Cell and Molecular Biology</u> (1987) Academic Press, London. 325 pp. (ISBN 0-12480-855-7).

Sambrook, J. et al. eds., Molecular Cloning: A Laboratory Manual (2d Ed. 1989) Cold Spring Harbor Laboratory Press, NY. Vols. 1-3. (ISBN 0-87969-309-6).

Winnacker, E.L. From Genes To Clones: Introduction To Gene Technology (1987) VCH Publishers, NY (translated by Horst Ibelgaufts). 634 pp. (ISBN 0-89573-614-4).

[0071] The present invention further incorporates by reference in their entirety techniques well known in the field of microarray construction and analysis. These techniques include, but are not limited to, techniques described in the following patents and patent applications describing arrays of biopolymeric compounds and methods for their fabrication: U.S. Pat. Nos. 5,242,974; 5,384,261; 5,405,783; 5,412,087; 5,424,186; 5,429,807; 5,436,327; 5,445,934; 5,472,672; 5,527,681; 5,529,756; 5,545,531; 5,554,501; 5,556,752; 5,561,071; 5,599,895; 5,624,711; 5,639,603; 5,658,734; 5,807,522; 6,087,102; WO 93/17126; WO 95/11995; WO 95/35505; EP 742 287; and EP 799 897. Techniques also include, but are not limited to, techniques described in the following patents and patent application describing methods of using arrays in various applications: U.S. Pat. Nos. 5,143,854; 5,288,644; 5,324,633; 5,432,049; 5,470,710; 5,492,806; 5,503,980; 5,510,270; 5,525,464; 5,547,839; 5,580,732; 5,661,028; 5,994,076; 6,033,860; 6,040,140; WO 95/21265; WO 96/31622; WO 97/10365; WO 97/27317; EP 373 203; and EP 785 280.

[0072] It is readily apparent to those skilled in the art that other suitable modifications and adaptations of the com-

positions and methods of the invention described herein are obvious and may be made without departing from the scope of the invention or the embodiments disclosed herein. Having now described the present invention in detail, the same will be more clearly understood by reference to the following examples, which are included for purposes of illustration only and are not intended to be limiting of the invention.

Examples

Materials and Methods

10 EXAMPLE 1:

Signal Transduction cDNA Microarray Construction

[0073] To study the cellular pathology associated HCMV, a unique microarray of more than 1100 signal transduction cDNAs was created.

[0074] In order to manufacture cDNA-arrays on membranes, the following strategy was employed: cDNAs encoding parts of or full length proteins of interest (referred to as "target cDNAs") were cloned into the plasmid BLUESCRIPT II KS+ (Stratagene, USA). Large scale purifications of these plasmids were performed according to standard techniques, and 200 μ l aliquots (1 μ g/ μ l plasmid concentration) were transferred into appropriate 96 well plates. The plates were then sealed with sealing tape, incubated for 10 minutes at 95°C, and chilled on ice for 5 minutes. 10 μ l of 0.6N NaOH were added, and the mix was then stored for 20 minutes at room temperature. Following the incubation at room temperature, 10 μ l 2.5M Tris-HCI (Tris-(hydroxymethyl)-aminomethane-hydrochloride) pH7.1 and 20 μ l 40x SSC (6M Sodium chloride - 0.6M tri-Sodium citrate buffer) was added.

[0075] Target cDNAs were spotted onto Nylon or Nitrocellulose membranes using a BIOGRID (BioRobotics, UK) equipped with a 0.7mm pintool. 200-350ng of plasmid-encoding target cDNAs were transferred onto the membranes and crosslinked to the membranes using ultraviolet light (1.2x10 5 μ J/cm 2) treatment. The arrays were stored for use in subsequent experiments (described below) at room temperature.

EXAMPLE 2:

30

35

40

45

50

55

HCMV Infection

[0076] To examine the effects of HCMV infection on cellular signal transduction activity, HCMV-infected cells were generated for comparison to control (i.e., uninfected) cells.

[0077] Primary human foreskin fibroblasts (HFF) were grown close to confluency in MEM medium (Minimum Essential Medium, Life Technologies) supplemented with 20% fetal calf serum at 37°C and 5% CO_2 to obtain $\sim 6x10^6$ cells per tissue culture flask. Virus adsorption to the cells was performed with the HCMV strain AD169 at different (0.2, 1, and 3) multiplicities of infection (MOI) for 90 minutes in a volume of 5ml at 37°C. The viral inoculum was removed, and cells were cultured in 50ml of MEM medium supplemented with 20% fetal calf serum and 150 μ g/ml cycloheximid at 37°C and 5% CO_2 for 7, 24, 48, or 72 hours, respectively.

EXAMPLE 3:

Isolation and Purification of Poly A+ RNA

[0078] In order to perform differential expression analysis using the cDNA microarray described in Example 1, RNA extraction and purification on the HCMV-infected and uninfected cells was performed using techniques known in the art. [0079] Briefly, after incubation (for the respective time-intervals) of infected and control cells, cells were washed twice with phosphate buffered saline (PBS) and then trypsinized. Cells were removed from the culture dish by resuspension with PBS. Cells were then sedimented, and directly lysed by repetitive pipetting in 1ml of Tri reagent (Molecular Research Centre, Inc., USA) per 1×10⁶ cells.

[0080] Cell lysates were stored at room temperature for 5 minutes, and then centrifuged (12,000xg) for 15 minutes at 4°C. The supernatant was mixed with 0.1ml of 1-bromo-3-chloropropane per 1ml of Tri reagent and shaken vigorously. The resultant suspension was stored for 5 minutes at room temperature, and then centrifuged (12,000xg) for 15 minutes at 4°C.

[0081] Following centrifugation, the colorless upper phase was transferred into new tubes, mixed with 5μ l of polyacryl-carrier (Molecular Research Centre, Inc., USA), and vigorously shaken with 0.5ml of isopropanol per 1ml of Tri reagent. The samples were stored at room temperature for 5 minutes and then centrifuged (12,000xg) for 8 minutes

at 4°C. The supernatant was removed and the RNA pellet washed twice with 1ml of 75% ethanol. The pellet was dried and resuspended in RNase-free buffer at a concentration of 1µg RNA per 1µl buffer.

[0082] Purification of poly A+ RNA from total RNA was performed using the OLIGOTEX system (Qiagen, Germany) following manufacturer's instructions. In brief, 100-200μg of total RNA was brought up to 250μl with RNase-free water, and 250μl of buffer OBB (20mM Tris/HCl pH7.5, 1M NaCl, 2mM EDTA, 0.2% SDS) and 15μl of OLIGOTEX suspension was added. The samples were incubated for 3 minutes at 70°C, and placed at room temperature for 10 minutes. The samples were centrifuged for 2 minutes (12,000xg), and the supernatant removed. The remaining pellet was resuspended in 400μl buffer OW2 (10mM Tris/HCl pH7.5, 150mM NaCl, 1mM EDTA). The suspension was transferred to a spin column (supplied with the system) and centrifuged at 12000xg for 1 minute at room temperature. The spin column was transferred to a new tube and 400 μl of buffer OW2 was applied on the column. The spin column was centrifuged (12,000xg) for 1 minute at room temperature. The spin column was transferred to a new tube and the RNA eluted from the column by the addition of 50μl buffer OEB (5mM Tris/HCl pH7.5) (at 70°C) to the column, resuspension of the Oligotex-resin, and centrifugation (12,000xg) for 1 minute at room temperature.

[0083] Any genomic DNA contamination of the RNA preparations was eliminated by enzymatic digestion using DNase I. 6 μ I of 10x DNase buffer (Promega, USA) and 4 μ I of RQ-DNase (Promega, USA) were added to 50 μ I of the RNA-buffer solution, and the reaction mixture was incubated for 15 minutes at 37°C. Stop-buffer (6 μ I; Promega, USA) was then added, the mixture brought to 200 μ I final volume with TE buffer (10mM Tris/HCI, 1mM EDTA), and Phenol/Chloroform extractions were performed twice. The RNA-containing phase was transferred to new reaction tubes and RNA was precipitated using 5M NaCI (final concentration of 0.2M), 1 μ I poly-acryl-carrier (Molecular Research Centre, Inc., USA) and 500 μ I of 100% ethanol. The solution was centrifuged for 10 minutes at 4°C, the RNA pellet washed with 1mI of 80% ethanol, dried, and resuspended in 30 μ I TE buffer. Poly A+ RNA suspension samples were stored at -70°C for use in subsequent experiments.

EXAMPLE 4:

10

20

25

30

35

40

45

50

Preparation of Radioactively Labeled cDNA Probes from RNA

[0084] To prepare test and control samples for microarray analysis, RNA samples isolated and purified from HCMV-infected and control cells (prepared as described in Example 3) were used to generate radioactively labeled cDNA probe. Many techniques to generate labeled cDNA constructs from cellular RNA extracts are known in the art and applicable to the present invention. Two of those protocols were used in this example to generate radiolabeled cDNA from RNA samples: the first technique involved reverse transcribing cDNA from the RNA sample in the presence of radioactively labelled dATP; the second technique involved first strand cDNA synthesis from the RNA sample, followed by random priming with radioactively labelled dATP.

[0085] For reverse transcription of cDNA from the RNA sample in the presence of radioactively labelled dATP, $1\mu g$ of primer TXN (5'-TTT TTT TTT TTT TTT TXN-3'; SEQ ID NO:1; with T = dTTP; N = dATP, dCTP, dGTP or dTTP; X = dATP, dCTP or dGTP) and total RNA (1 to 15 μg) or poly A+ RNA (20 to 500 ng) were combined in 12 μl bidistilled DEPC-treated H₂O (DEPC: diethylpyrocarbonate) and shaken for 5-15 minutes at 60°C. The mixture was then incubated at 4°C for 2-10 minutes, and centrifuged (10,000xg) for 30 seconds.

[0086] After centrifugation, 7μl of a labelling mix (100μCi χ[33P]-ATP (Amersham, UK); vacuum dried and resuspended in 4μl first strand buffer (Life Technologies, USA); 2μl 0.1M DTT (dithiothreitol); and 1μl labelling solution -- 4mM dCTP, dGTP, dTTP each and 80 μM dATP final concentration) was added to the RNA solution. 1μl SUPERSCRIPT II reverse transcriptase (Life Technologies, USA) was added and the reaction incubated for 10 minutes at room temperature and then for 60 minutes at 38°C. Following the reaction incubation, 5μl 0.5M EDTA (ethylene diamine tetraacetate) and 25 μl 0.6M NaOH was added to the reaction mixture and shaken vigorously for 30 minutes at 68°C.

[0087] Unincorporated nucleotides were removed from the labelling reaction using PROBEQUANT G-50 columns (Amersham, UK). The column (with bottom closure and lid removed) was shaken vigorously and centrifuged (735xg) for 1 minute in an appropriate reaction tube. The column was placed into a new reaction tube, the probe was applied onto the center of the column material and the column was centrifuged (735xg) for 2 minutes. The flow-through was transferred into new reaction tubes and bidistilled H₂O added to 100 µl final volume. 5M NaCl, 1µl poly-acryl-carrier (Molecular Research Centre, Inc., USA) and 250 µl ethanol was added, and the probe precipitated by centrifugation (12,000xg) for 15 minutes. The supernatant was discarded and the pellet dried for subsequent use.

[0088] For the alternate labelling technique (random priming with radioactively labelled dATP after first strand cDNA synthesis), the following procedure was followed: 1μg primer TXN (see above) was added to 20-500ng of poly A+ RNA in 12 μl final volume, incubated for 5 minutes at 60°C, followed by an addition incubation for 2-10 minutes on ice. The mix was centrifuged (12,000xg) for 30 seconds, and 4μl of first strand buffer (Life Technologies, USA), 2μl 0.1M DTT, 1μl 10mM dNTP and 1μl SUPERSCRIPT II reverse transcriptase (Life Technologies, USA) was added. The reaction was incubated for 10 minutes at room temperature, followed by an additional incubation for 60 minutes at 38°C. Fol-

lowing the reaction incubation, 5μ I 0.5M EDTA and 25 μ I 0.6M NaOH was added to the reaction mixture and shaken vigorously for 30 minutes at 68°C.

[0089] Unincorporated nucleotides were removed as described above; however, the final pellet was resuspended in 30μ l bidistilled H_2O .

[0090] 15 μ l of the resuspended cDNA solution was transferred to new reaction tubes, incubated for 5 minutes at 95°C, chilled on ice for 5 minutes, and centrifuged for 30 seconds. Following manufacturer's instructions accompanying the Random Primers DNA Labelling system (Life technologies, USA), 15 μ l buffers mixture, 2 μ l of each dCTP, dGTP and dTTP (provided with the system) were added to the cDNA. 5 μ l γ [33P]-ATP (Amersham, UK) was added and the mixture adjusted to 49 μ l final volume with bidistilled H₂O. The reaction was started by addition of 1 μ l Klenow enzyme (supplied with the system), and incubated for 60 minutes at 25°C 5 μ l. Stop solution (provided with the system) was added and unincorporated nucleotides were removed by column purification as described above.

EXAMPLE 5:

10

15 Hybridization of Labeled cDNA Probe to cDNA Array

[0091] To screen HCMV-infected cells compared to uninfected cells for differential activation of cellular signal transduction, labeled cDNA probes (generated according to Example 4) were exposed to a signal transduction cDNA microarray (generated as described in Example 1) following hybridization techniques known in the art.

[0092] Sample pellets from Example 4 were resuspended in 10μ I C_0 T DNA $(1\mu g/\mu I)$, Roche Diagnostics, Germany), 10μ I yeast tRNA $(1\mu g/\mu I)$ Sigma, USA) and 10μ I poly A $(1\mu g/\mu I)$, Roche Diagnostics, Germany). Herring sperm DNA (to a final concentration of $100\mu g/mI$), 5μ I 10% SDS (Sodiumdodecylsulfate), and 25μ I 20x SSPE was added, and adjusted 100μ I final volume with bidistilled H_2 O. The mix was incubated for 5 minutes at 95° C, centrifuged (10,000xg) for 30 seconds, and vigorously shaken for 60 minutes at 68° C. A 1μ I aliquot of the probe was used to measure the incorporation of radioactive dATP with a scintillation counter. Probes with at least a total of $20x10^{6}$ cpm were used for the screen assay.

[0093] Arrays were prehybridized in hybridization solution for at least 30 minutes in a roller bottle oven at 42°C. Following prehybridization, radiolabelled probe was added to the hybridization solution and hybridization was continued for 20-40 hours.

[0094] Following hybridization, the probe was discarded and the array subjected to a series of washes. Initially the arrays were washed twice in wash solution A (2xSSC) in the roller oven at room temperature. Wash solution A was then replaced with wash solution B (2x SSC, 0.5% SDS), preheated to 60°C, and arrays were washed twice for 30 minutes at 60°C. Wash solution B was then replaced with wash solution C (0.5x SSC, 0.5% SDS), preheated to 60°C, and arrays were washed twice for 30 minutes at 60°C.

35 [0095] The moist arrays were wrapped in airtight bags and exposed for 8-72 hours on erased phosphoimager screens (Fujifilm, Japan).

EXAMPLE 6:

50

55

40 Signal Transduction cDNA Array Analysis

[0096] To demonstrate differential activation of cellular signal transduction in HCMV-infected cells compared to uninfected cells, hybridized cDNA arrays from infected and uninfected samples were analyzed.

[0097] Exposed phosphoimager screens (from Example 5) were scanned with a resolution of 100μ and 16bits per pixel using a BAS-1800 (Fujifilm, Japan). The data were imported into the computer program, ARRAYVISION (Imaging Research, Canada), and analyzed according the computer program's specification. Hybridization signal strength is indicative of the quantity of RNA molecules present in the probe. Differentially expressed genes were identified according to the ratio of signal strength after normalization to the overall intensity of the arrays.

[0098] Signal transduction cDNA microarray analysis of radiolabelled cDNA-probes from HCMV-infected (strain AD169) versus non-infected primary human foreskin fibroblasts to cDNA-arrays revealed significant upregulation of the cellular kinase cDNAs:

RICK (2-fold at 3 hours post infection; 3.6-fold at 7 hours post infection);

RIP (2.6-fold at 3 hour post infection; 2.2-fold at 24 hour post infection);

NIK (4-fold at 7 hour post infection);

MKK3 (2-fold at 3 hour post infection; 2.5-fold at 7 hour post infection); and

SRPK-2 (2.2-fold at 7 hour post infection)

compared to uninfected human foreskin fibroblasts cells.

EXAMPLE 7:

10

Northern Blot Analysis

[0099] To confirm the results of the microarray analysis of Example 6, northern blot analysis was performed according to techniques well known in the art.

[0100] HCMV-infected and uninfected cells (from Example 2) cells were pelleted and the total RNA was prepared as follows: Following centrifugation and removal of the supernatant, cells were lysed in 1ml of Trizol reagent (readyto-use-reagent from Gibco-BRL) per 1.5 106 cells. The Tri reagent/cell lysate was transferred to an eppendorff tube and centrifuged (13,000rpm) for 15 minutes at 4°C. The supernatant was transferred to a new eppendorff tube and 0.1ml of BCP (1-bromo-3-chlorpropane) for each ml of Tri reagent was added. Samples were vortexed for 15 seconds, incubated for 5 min at room temperature, and then centrifuged (13,000rpm) for 15 minutes at 4°C. The upper aqueous phase was transferred to a new eppendorff tube, 0.5 ml isopropanol was added for each ml of Tri reagent (Molecular Research Center, Inc., USA), vortexed, and incubated for additional 8 min at room temperature, and centrifuged (13,000rpm) for 10 min at 4°C. The supernatant was aspirated, and the precipitated RNA was washed twice with icecold 75% ethanol and air-dried. The RNA pellet was resuspended in 50µl Tris-HCl pH 7.5.

[0101] The quantity of the RNA for each sample was determined by UV-spectroscopy, and the quality was determined via gel electrophoresis on a formaldehyde-containing 1.2% agarose gel.

[0102] RNA samples of 10µg each were size-fractionated by 1.2% formaldehyde agarose gel electrophoresis and transferred to synthetic membrane filters (Hybond N, Amersham) with 20XSSC (1 X SSC is 150 mM NaCl, 15 mM C₆H₅Na₃O₇ x 2H₂O, pH 7.0) overnight. RNA was immobilized to the filter using UV-light for crosslinking (120 mJ/cm² for 25 seconds).

[0103] Membrane filters were firstly prehybridized for 4 hours at 65°C in a prehybridization solution containing 5 X SSC, 10 X Denhardt's solution (1 X Denhardt's solution is 0.02% bovine serum albumine, 0.02% polyvinyl pyrrolidone, 0.02% ficoll), 20mM sodium phosphate, pH 7.0, 7% SDS, 100μg/ml sonicated salmon sperm DNA, and 100μg/ml. Hybridization was performed at 65°C in the prehybridization buffer containing 10% dextran sulphate, plus added radiolabelled probe for 16 hours.

30 [0104] Membrane filters were hybridized to oligonucleotide probes specific for a particular cellular kinase identified in Example 6. Probes sequences included the following:

Table 4:

Cellu	ılar	cDNA Probe Sequence	SEQ
Kina	se		ID NO:
NI	< 5'- GTC C	TG GAG GGC TCT TTT TGA TGA AAC C – 3'	2
RI	5'- GTG C	TC AAT GCA GTT GGG CCC CTT GTA CAC-3'	3
RIC	K 5'- GTC G	AG CAG CGG AGT GTG GAT GTG CAG – 3'	4

[0105] The oligonucleotides were radiolabelled at their 3' ends with (alpha-32P) deoxyadenosine 5'- triphosphate $(^{32}P-\alpha-dATP)$ (Amersham) employing the Terminal Transferase kit (Roche) following manufacturer's instructions.

[0106] Unincorporated ³²P-α-dATP nucleotides were removed similar to the protocol described in Example 4: After vortexing the PROBEQUANT Sephadex G-50 (Amersham, UK) column (with bottom closure and lid removed), the column was placed in a 2 ml tube and centrifuged for 1 minute at 735xg. The column was placed in a new 1.5ml eppendorff tube (without a cap), and the radioactive probe was pipetted carefully on the center of the preformed resin. Centrifugation (735xq) for 2 minutes effectively removed the unincorporated ³²P-α-dATP nucleotides.

[0107] Hybridized filters were washed once in 5% SDS, 3 X SSC, 10 X Denhardt's solution, 20mM sodium phosphate, pH 7.0 for 30 min at 65°C. A second wash step followed in 1 X SSC, 1% SDS at 65°C for 30 min.

[0108] Filters were exposed at -80°C to Kodak XAR-5 films using intensifying screens.

[0109] Northern blot analysis confirmed upregulation of cellular kinase mRNA: RICK, RIP, and NIK in HCMV-infected cells compared to uninfected cells, consistent with results obtained from microarray analysis.

15

35

40

45

50

EXAMPLE 8:

10

25

30

35

40

50

Western Blot Analysis

[0110] To further confirm the results of the microarray analysis of Example 6 and northern blot analysis of Example 7, western blot analysis was performed according to techniques well known in the art.

[0111] HCMV-infected and uninfected cells (from Example 2) were pelleted and polypeptide extracts prepared as follows: Infected and uninfected cell samples (from various time intervals) were lysed with 420 μ l of lysis buffer (20mM Hepes (N-[2-hydroxyethyl]piperazine-N'-[2-ethansulfonic acid]) pH7.5, 150 mM NaCl, 1% TRITON X-100 (t-octylphenoxypolyethoxyethanol), 10% glycerol, 1mM PMSF (phenylmethylsulfonyl fluoride), 10 μ g/ml Aprotinin, 1mM orthovanadat) on ice. Lysed cells were cleared from debris by centrifugation (15 minutes, 13000 rpm, 4°C), dissolved in 1x Laemmli buffer, denatured for 5 minutes at 100°C and submitted to SDS-PAGE (gradient gel 7% - 12%).

[0112] Gels were blotted onto nitrocellulose filters (Amersham, UK) for 3 hours (0.8mA/cm2). Detection of expression of the identified host cell kinases was performed using the following target specific antibodies: OPA1-01023 polyclonal rabbit anti-RICK antibody (Dianova); H-207 polyclonal rabbit anti-RIP antibody (Santa Cruz Biotechnology); I-20 polyclonal rabbit anti-MKK3 antibody (Santa Cruz Biotechnology); S80620 murine anti-SRPK2 antibody (Transduction Laboratories); anti-NIK rabbit serum (generated by SIGMA Genosys Biotechnologies using the NIK-peptide 5'-CNPT-NTRPQSDTPEIRKYKKRFN-3', SEQ ID NO:5, for immunization). All antibodies were used according to the manufacturer's instructions. Detections were performed with the ECL Kit (Amersham, UK).

[0113] Western blot analysis confirmed the transcriptional upregulation of infected host cell kinase mRNAs resulted in increased expression of the respective proteins:

A single ~60kDa band representing RICK was upregulated between 7-24 hours post HCMV infection;

A single ~74kDa band representing RIP was upregulated between 7-72 hours post HCMV infection;

A single ~135kDa band representing NIK was upregulated between 24-72 hours post HCMV infection;

A single ~35kDa band representing MKK3 was upregulated between 7-72 hours post HCMV infection; and

A single \sim 115kDa band representing SRPK2 was upregulated between 24-72 hours post HCMV infection.

EXAMPLE 9:

Genetic Validation

[0114] HFF cells were infected with Adenovirus expressing various kinase constructs at different particles per cell ratios (p/c). The adenovirus used here were all E1, E3 defective derivatives of adenovirus type 5 (reviewed in Russell WC (2000) Update on adenovirus and its vectors. J Gen Virol. 81:2573-604). Briefly, the cDNA of interest was cloned into a transfer plasmid bearing the CMV IE promoter enhancer (IE: immediate early) and the rabbit beta-globin intron/polyadenylation signal. This expression cassette was inserted into a bacterial plasmid borne-adenovirus genome using recombination in bacteria (Chartier C., E. Degryse, M. Gantzer, A. Dieterle, A. Pavirani, and M. Mehtali. 1996. Efficient generation of recombinant adenovirus vectors by homologous recombination in *Escherichia coli*. J. Virol. 70: 4805-4810.). Virus was amplified in HEK 293 cells and purified from cell lysates using CsCl density gradient centrifugation as described (Cotten, M., Baker A., Birnstiel M.L., Zatloukal, K., Wagner, E. (1996) Adenovirus polylysine DNA conjugates. in Current Protocols in Human Genetics, Eds. N. C. Dracopoli, J. L. Haines, B.R. Korf, D.T. Moir, C.C. Morton, C.E. Seidman, J.G. Seidman, D.R. Smith; John Wiley and Sons, Inc. New York. pp. 12.3.1-12.3.33.). The control viruses AdJ5 was previously described (Glotzer J.B., Saltik M., Chiocca S., Michou A.I., Moseley P. and Cotten M. (2000) Activation of heat-shock response by an adenovirus is essential for virus replication. Nature 407:207-11).

[0115] Two days after plating HFF cells, cultures were infected with CMV strain Ad169-GFP. Replication of CMV was estimated after one week (7 dpi) utilizing the GFP-signal expressed as GFP counts.

[0116] Fig. 1 shows the reduction rates in HCMV replication of HHF cells pre-infected with Adeno virus containing the RIP wildtype sequence (AdRIPwt; 1 - 4) and a RIP inactive mutant (AdRIPkr; 5 - 8).

[0117] No HCMV-infection resulted in hardly any signal (mock, 13), while infection with HCMV yielded in about 13.000 GFP counts (AD169-GFP; 14). Pre-infection with increasing amounts of control Adeno virus (AdliteJ5) caused a slight reduction in HCMV replication (9 - 12). There was a clear difference, when HFF cells were pre-infected with Adeno virus containing the RIP wildtype sequence (AdRIPwt; 1 - 4) and a RIP inactive mutant (AdRIPkr; 5 - 8). The lysine (K) at amino acid position 45 is mutated to an arginine (R), which renders the kinase inactive. This mutation was introduced into the human RIP cDNA utilizing the QuikChangeTM Site-directed Mutagenesis Kit (Stratagene, CA, USA) according to the instructions of the manufacturer. Expression of the mutated RIP kinase efficiently blocked HCMV replication (5 - 8), while the wildtype sequence was less potent in doing so (1 - 4).

[0118] Fig. 2 shows the reduction rates in HCMV replication of HHF cells pre-infected with Adeno virus containing

the RICK wildtype sequence (AdRICKwt) and two RICK inactive mutants (AdRICKkr and AdRICKdn). In one construct (AdRICKkr), the lysine (K) at position 47 is mutated to an arginine (R). In the other construct (AdRICKdn), the aspartate at position 146 is mutated to an asparagine. Both changes in sequence render the kinase inactive. The mutations were introduced into the human RICK cDNA utilizing the QuikChange™ Site-directed Mutagenesis Kit (Stratagene, CA, USA) according to the instructions of the manufacturer.

[0119] Similar experiments as described for RIP (Example 9, Fig. 1) were also performed with RICK. No HCMV-infection resulted in hardly any signal (Mock + Mock), while infection with HCMV yielded in about 7.000 GFP counts (Mock + AD169-GFP). Pre-infection with increasing amounts of control Adeno virus (AdliteJ5, from 111 to 3000 particles per cell) caused a slight reduction in HCMV replication. There was a clear difference, when HFF cells were pre-infected with Adeno virus containing the RICK wildtype sequence (AdRICKwt and two RICK inactive mutants (AdRICKkr and AdRICKdn). All three RICK constructs efficiently reduced HCMV-replication.

EXAMPLE 10:

10

30

40

45

50

55

15 RICK-Kinase Assay

[0120] To obtain active RICK kinase the human RICK-cDNA was fused with a DNA sequence coding for the HA-tag and cloned into the eucaryotic expression vector pcDNA3 (Invitrogene). This construct was transfected into human embryonic kidney cells (HEK 293) via the calcium-phosphate co-precipitation method. One day after transfection medium was replaced by fetal calf serurum-free medium and two days after transfection cells were washed with PBS and harvested and lysed in RIPA-buffer (150 mM NaCl, 1 mM EDTA, 1% Tritron X-100, 1% Na-desoxycholate, 0.1 % SDS, 10 mM Tris-HCl pH 7.5). The RICK-HA fusion protein was immunoprecipitated from 250 μl cleared lysate (i.e. lysate of one well of a six-well plate) utilizing an anti-HA antibody from Roche Pharmaceuticals and Protein A sepharose. After addition of 500 μl of HNTG-buffer (50 mM HEPES pH 7.5, 150 mM NaCl, 1 mM EDTA, 10% glycerine, 0.1% Triton X-100) the sample was rotated for 3 hrs at 4°C. After washing the immunoprecipitate twice with 0.5 ml HNTG-buffer and twice with 0.5 ml assay-buffer (25 mM Tris-HCl pH 7.5, 3 mM MgCl₂, 0.5 mM MnCl₂, 0.5 mM DTT and 50 mM NaCl), the kinase reaction was performed directly on the beads in 40 μl assay buffer containing 2.5 μCi γ [33P]-ATP and various concentrations (between 100 nM and 50 μM) of compounds of Table 1. After 30 min at 30°C, the reaction was stopped by addition of 40 μl 3X Laemmli-buffer (16% glycerol, 1.01M β-mercaptoethanol, 5% SDS, 200mM Tris/HCl pH 6.8, 8% bromphenolblue). Phosphorylation products were analyzed by SDS-PAGE and auroradiography (x-ray film and phosphor imager).

EXAMPLE 11:

35 RIP-Kinase Assay

[0121] To obtain active RIP kinase the human RIP-cDNA was fused with a DNA sequence coding for the HA-tag and cloned into the eucaryotic expression vector pcDNA3 (Invitrogene). This construct was transfected into human embryonic kidney cells (HEK 293) via the calcium-phosphate DNA co-precipitation method. Two days after transfection cells were washed with PBS and harvested and lysed in lysis-buffer (150 mM NaCl, 1 mM EDTA, 1% Tritron X-100, 20 mM Tris-HCl pH 7.5 and freshly added: 30 mM NaF, 10 μg/ml Aprotinine, 10 μg/ml Leupeptine, 2 mM Na-pyrophosphate). The RIP-HA fusion protein was immunoprecipitated from 250 μl cleared lysate (i.e. lysate of one well of a six-well plate) utilizing an anti-HA antibody from Roche Pharmaceuticals and Protein A sepharose. The sample was rotated for 3 hrs at 4°C. The immunoprecipitates were washed twice with 0.75 ml lysis-buffer, twice with 0.75 ml high salt-buffer (1 M NaCl, 1 mM EDTA, 1% Tritron X-100, 20 mM Tris-HCl pH 7.5 and freshly added: 30 mM NaF, 10 μg/ml Aprotinine, 10 μg/ml Leupeptine, 2 mM Na-pyrophosphate), twice with 0.75 ml lysis-buffer and twice 0.75 ml with kinase assay buffer (10 mM MgCl₂, 10 mM MnCl₂, 10 mM benzamidine, 0.5 mM EDTA). The kinase reaction was performed directly on the beads in 40 μl kinase assay buffer containing 2.5 μCi γ [3²P]-ATP and various concentrations (between 100 nM and 50 μM) of compounds of Table 2. After 30 min at 30°C, the reaction was stopped by addition of 40 μl 2X Laemmli-buffer. Phosphorylation products were analyzed by SDS-PAGE and autoradiography (x-ray film and phosphor imager).

EXAMPLE 12:

Virus Replication Assay

Cell culture and virus

[0122] Primary human foreskin fibroblasts (HFF) were cultivated in MEM containing 5% (v/v) fetal calf serum. Infec-

tion analysis was restricted to cell passage numbers below twenty. Human cytomegalovirus strain AD169 (ATCC) was grown in HFF cells and quantitated for infectivity by the plaque reduction assay. Aliquots were stored at -80°C.

Construction of recombinant cytomegalovirus

[0123] For construction of a recombination vector, two linker sequences were inserted into the pBlueScribe vector pBS+ (Stratagene): the first contained restriction sites for Nhel, Spel, Pacl and Bgll followed by a loxP sequence (ATAACTTCGTATAGCATACATTATACGAAGTTAT) and was introduced into Pstl/Xbal sites of the vector; the second contained another loxP sequence followed by restriction sites Hpal, Clal and Pmel and was introduced into BamHI/ Asp718 sites. A gene cassette comprising of a "humanized" version of the ORF coding for GFP (gfp-h) under the control of the HCMV enhancer/promoter and the Ptk/PY441 enhancer-driven neoR selection marker was excised from plasmid pUF5 (Zolotukhin et al., 1996, J. Virol. 70, 4646-4654) and inserted into the recombination vector via BgIll sites. [0124] At the 5' and 3'-positions of this loxP-flanked gene cassette, two HCMV sequences with homology to the gene region containing the open reading frames US9 and US10 were inserted. For this, viral sequences were amplified from template pCM49 (Fleckenstein et al., 1982, Gene 18, 39-46) via PCR in a 35-cycle program (denaturation 45 sec at 95°C, annealing 45 sec at 55°C and elongation 2 min at 72°C) by the use of Vent DNA polymerase (New England Biolabs). A US10-specific sequence of 1983 bp in length was generated using primers US10[200900]Spel (GCT-CACTAGTGGCCTAGCCTGGCTCATGGCC) and US10[198918]Pacl (GTCCTTAATTAAGACGTGGTTGTGGTCAC-CGAA) and inserted at the vector 5' cloning position via Spel/Pacl restriction sites (see bold-print). A US9-specific sequence of 2010 bp was generated using primers US9-3'Pmel (CTCGGTTTAAACGACGTGAGGCGCTCCGTCACC) and US-5' Clal (TTGCATCGATACGGTGTGAGATACCACGATG) inserted at the vector 3' cloning position via Pmel/ Clal restriction sites.

[0125] The resulting construct pHM673 was linearized by the use of restriction enzyme Nhel and transfected into HEF cells via the electroporation method using a Gene Pulser (Bio-rad; 280 V, 960 μ F, 400 Ω). After 24 h of cultivation, cells were used for infection with 1 PFU/ml of HCMV strain AD169. Selection with 200 μ g/ml G418 was started 24 h post infection. Following 3 weeks of passage in the presence of G418, GFP fluorescence could be detected in most of the infected cells. Plaque assays were performed with infectuous culture supernatant on HFF cells and single virus plaques were grown by transfer to fresh HFF cells cultured in 48-well plates. DNA was isolated from cells of 32 fluorescence-positive wells and confirmed for the presence of recombinant virus by PCR. For this, primers US9[198789] (TGACGCGAGTATTACGTGTC) and US10[199100] (CTCCTCCTGATATGCGGTT) were used resulting in an amplification product of 312 bp for wild-type AD169 virus and approximately 3.5 kb for recombinant virus.

Plaque assay

10

30

35

40

50

55

[0126] HFF cells were cultivated in 12-well plates to 90-100% confluency and used for infection with dilutions of virus-positive cell culture supernatants. Virus inoculation was performed for 90 min at 37°C under occasional shaking before virus was removed and the cell layers were rinsed with PBS. Overlays of MEM 5% (v/v) fetal calf serum and 0.3% (w/v) agarose were added to each well and all samples were incubated at 37°C in a 5% CO₂ atmosphere for approximately 12 days. Finally, overlays were removed and the formation of foci was visualized by staining with 1 % crystal violet in 20% ethanol for 1 min. After repeated rinsing with PBS, plates were air-dried at room temperature and plaque numbers were counted with a light microscope. For the recombinant AD169-GFP virus, quantification of plaque assays could also be performed without crystal violet staining by a direct counting of the amount of green fluorescent plaques using fluorescence microscopy.

45 Antiviral compounds

[0127] The reference compounds used for antiviral studies, ganciclovir (GCV, Cymeven), foscarnet sodium (FOS, Foscavir) and cidofovir (CDV, Vistide) were purchased from Syntex Arzneimittel (Aachen, Germany), Sigma-Aldrich (Germany) and Pharmacia & Upjohn S.A. (Luxembourg), respectively. Stocks were prepared in aequeous solution and stored at -20°C. The test compounds were dissolved in DMSO and aliquots were stored at -20°C.

GFP infection assay

[0128] HFF cells were cultivated in 12-well plates to 90-100% confluency and used for infection with 0,5xTCID₅₀ of AD169-GFP virus. Virus inoculation was performed for 90 min at 37°C with occasional shaking before virus was removed and the cell layers were rinsed with PBS. Infected cell layers were incubated with 2 ml of MEM containing 5% (v/v) fetal calf serum and optionally of the respective test substances or DMSO as control. Infected cells were incubated at 37°C in a 5% CO₂ atmosphere for 7 days and harvested by trypsination and centrifugation. 200 µl of lysis buffer (25

mM Tris pH 7.8, 2 mM DTT, 2 mM trans-1,2-diaminocyclohexane-N,N,N',N'-tetraacetic acid, 1% Triton X-100, 10% glycerol) was added to each cell pellet and lysis was achieved by incubation for 10 min at 37°C followed by a 30-min incubation at room temperature on a shaker. Lysates were centrifuged for 5 min at 15.000 rpm in an Eppendorf centrifugue to remove cell debris. Supernatants were transferred to an opaque 96-well plate for automated measuring of GFP signals in a Victor 1420 Multilabel Counter (Wallac). GFP units were converted to percent inhibition values relative to DMSO controls (set at 100% GFP expression).

Indirect immunofluorescence analysis

[0129] Cells were either grown on Lab-Tek Permanox slides (Nunc) or harvested from 6-well plates, spotted onto glass slides with marked rings (Medco) and fixed by a 15-min treatment with 3% formaldehyde in PBS followed by permeabilization for 15 min in 0.1% Triton X-100 in PBS at room temperature. Blocking was achieved by incubation with Cohn Fraction II/III of human gamma-globulin (Sigma; 2 mg/ml) for 30 min at 37°C. The IE1/IE2-specific primary antibody MAb810 (Chemicon International, Inc. CA, USA; dilution 1:10.000) was incubated for 90 min, the secondary antibody (tetramethyl rhodamine [TRITC]-coupled anti-mouse antibody, Dianova, dilution 1:100) for 45 min at 37°C before analysis by fluorescence microscopy. In addition to indirect TRITC staining of IE1/IE2 proteins, GFP signals could be detected directly via the fluorescence isothiocyanate (FITC) channel. Nuclear counterstaining was carried out using Vectashield mounting medium including DAPI (Vector Laboratories, Burlingame, CA).

SEQUENCE LISTING

5	<110> Axxima Pharmaceuticals AG
	<120> CELLULAR KINASES INVOLVED IN CYTOMEGALOVIRUS INFECTION AND THEIR INHIBITION
	<130> PA012EP
10	<140> <141>
	<160> 2
15	<170> PatentIn Ver. 2.1
	<210> 1 <211> 2501 <212> DNA <213> Homo sapiens
20	<220> <221> CDS <222> (225)(1847)
	<pre><400> 1 qqcaccaqtc tctaqaaaaq aagtcagctc tggttcggag aagcagcggc tggcgtgggc 60</pre>
25	cateegggga atgggegeec tegtgaeeta gtgttgeggg gcaaaaaggg tettgeegge 120
	ctcgctcgtg caggggcgta tctgggcgcc tgagcgcgca gtgggagcct tgggagccgc 180
30	cgcagcaggg ggcacacccg gaaccggcct gagcgcccgg gacc atg aac ggg gag 236 Met Asn Gly Glu
35	gcc atc tgc agc gcc ctg ccc acc att ccc tac cac aaa ctc gcc gac Ala Ile Cys Ser Ala Leu Pro Thr Ile Pro Tyr His Lys Leu Ala Asp 5 10 15 20
	ctg cgc tac ctg age cgc ggc gcc tct ggc act gtg tcg tcc gcc cgc Leu Arg Tyr Leu Ser Arg Gly Ala Ser Gly Thr Val Ser Ser Ala Arg 25 30 35
40	cac gca gac tgg cgc gtc cag gtg gcc gtg aag cac ctg cac atc cac His Ala Asp Trp Arg Val Gln Val Ala Val Lys His Leu His Ile His 40 45 50
45	act ccg ctg ctc gac agt gaa aga aag gat gtc tta aga gaa gct gaa 428 Thr Pro Leu Leu Asp Ser Glu Arg Lys Asp Val Leu Arg Glu Ala Glu 55 60 65
	att tta cac aaa gct aga ttt agt tac att ctt cca att ttg gga att Ile Leu His Lys Ala Arg Phe Ser Tyr Ile Leu Pro Ile Leu Gly Ile 70 75 80
50	tgc aat gag cct gaa ttt ttg gga ata gtt act gaa tac atg cca aat Cys Asn Glu Pro Glu Phe Leu Gly Ile Val Thr Glu Tyr Met Pro Asn 85 90 95 100
55	gga tca tta aat gaa ctc cta cat agg aaa act gaa tat cct gat gtt 572 Gly Ser Leu Asn Glu Leu Leu His Arg Lys Thr Glu Tyr Pro Asp Val 105 110 115

_	gct Ala	tgg Trp	cca Pro	ttg Leu 120	aga Arg	ttt Phe	cgc Arg	atc Ile	ctg Leu 125	cat His	gaa Glu	att Ile	gcc Ala	ctt Leu 130	ggt Gly	gta Val	620
5	aat Asn	tac Tyr	ctg Leu 135	cac His	aat Asn	atg Met	act Thr	cct Pro 140	cct Pro	tta Leu	ctt Leu	cat His	cat His 145	gac Asp	ttg Leu	aag Lys	668
10	act Thr	cag Gln 150	aat Asn	atc Ile	tta Leu	ttg Leu	gac Asp 155	aat Asn	gaa Glu	ttt Phe	cat His	gtt Val 160	aag Lys	att Ile	gca Ala	gat Asp	716
15	ttt Phe 165	ggt Gly	tta Leu	tca Ser	aag Lys	tgg Trp 170	cgc Arg	atg Met	atg Met	tcc Ser	ctc Leu 175	tca Ser	cag Gln	tca Ser	cga Arg	agt Ser 180	764
	agc Ser	aaa Lys	tct Ser	gca Ala	cca Pro 185	gaa Glu	gga Gly	Gly ggg	aca Thr	att Ile 190	atc Ile	tat Tyr	atg Met	cca Pro	cct Pro 195	gaa Glu	812
20	aac Asn	tat Tyr	gaa Glu	cct Pro 200	gga Gly	caa Gln	aaa Lys	tca Ser	agg Arg 205	gcc Ala	agt Ser	atc Ile	aag Lys	cac His 210	gat Asp	ata Ile	860
25		agc Ser															908
		gaa Glu 230															956
30		cat His															1004
35		cga Arg	_	-	_					_	_			_			1052
		gat Asp	-	-						_			_		-		1100
40		ttg Leu															1148
45		aag Lys 310			_		_	-	-		_	-				_	1196
		aag Lys															1244
50	Pro	caa Gln	Glu	Glu	Ser 345	Cys	Gly	Ser	Ser	Gln 350	Leu	His	Glu	Asn	Ser 355	Ğly	1292
55		cct Pro															1340

5	tta Leu	tct Ser	aga Arg 375	aaa Lys	gct Ala	caa Gln	gac Asp	tgt Cys 380	tat Tyr	ttt Phe	atg Met	aag Lys	ctg Leu 385	cat His	cac His	tgt Cys	1388
	cct Pro	gga Gly 390	aat Asn	cac His	agt Ser	tgg Trp	gat Asp 395	agc Ser	acc Thr	att Ile	tct Ser	ggt Gly 400	tct Ser	caa Gln	agg Arg	gct Ala	1436
10	gca Ala 405	ttc Phe	tgt Cys	gat Asp	cac His	aag Lys 410	acc Thr	act Thr	cca Pro	tgc Cys	tct Ser 415	tca Ser	gca Ala	ata Ile	ata Ile	aat Asn 420	1484
15	cca Pro	ctc Leu	tca Ser	act Thr	gca Ala 425	gga Gly	aac Asn	tca Ser	gaa Glu	cgt Arg 430	ctg Leu	cag Gln	cct Pro	ggt Gly	ata Ile 435	gcc Ala	1532
	cag Gln	cag Gln	tgg Trp	atc Ile 440	cag Gln	agc Ser	aaa Lys	agg Arg	gaa Glu 445	gac Asp	att Ile	gtg Val	aac Asn	caa Gln 450	atg Met	aca Thr	1580
20	gaa Glu	gcc Ala	tgc Cys 455	ctt Leu	aac Asn	cag Gln	tcg Ser	cta Leu 460	gat Asp	gcc Ala	ctt Leu	ctg Leu	tcc Ser 465	agg Arg	gac Asp	ttg Leu	1628
25	atc Ile	atg Met 470	aaa Lys	gag Glu	gac Asp	tat Tyr	gaa Glu 475	ctt Leu	gtt Val	agt Ser	acc Thr	aag Lys 480	cct Pro	aca Thr	agg Arg	acc Thr	1676
	tca Ser 485	aaa Lys	gtc Val	aga Arg	caa Gln	tta Leu 490	cta Leu	gac Asp	act Thr	act Thr	gac Asp 495	atc Ile	caa Gln	gga Gly	gaa Glu	gaa Glu 500	1724
30															atg Met 515		1772
<i>35</i>	ctt Leu	cag Gln	cct Pro	tac Tyr 520	ccg Pro	gaa Glu	ata Ile	ctt Leu	gtg Val 525	gtt Val	tct Ser	aga Arg	tca Ser	cca Pro 530	tct Ser	tta Leu	1820
					aat Asn				taa	gtga	actgt	tt t	tcaa	agaaq	ga		1867
40	aatg	gtgti	tc a	ataaa	aagga	at at	ttat	catc	ctç	gttgo	ettt	gact	tttt	tt a	atata	aaatc	1927
		-		_				-					_			atgaca	
																tacat	
45	_						_		_						-	tattc	
	_						_				_		_			cattca	
50	•		_			_						_				tgccc	
	taca	aagg	ggt t	tatta	aatt	a aa	aacto	ccatt	t att	agga	atta	catt	ttaa	aag 1	tt tt a	atttat	2347
	gaat	tcc	ett t	taaaa	aatga	at a	ttca	aaag	g taa	aaaca	aata	caat	ata	aag a	aaaa	aataa	2407
55	atat	atta	aat a	accg	gctto	ed to	gtcc	ccati	t tt t	caac	ctca	gcct	tcc	cta (ctgto	caccaa	2467

caaccaaqct aaataaagtc aacagcctga tgtg 2501

```
<210> 2
          <211> 540
          <212> PRT
          <213> Homo sapiens
          <400> 2
10
          Met Asn Gly Glu Ala Ile Cys Ser Ala Leu Pro Thr Ile Pro Tyr His
                                            10
           1
          Lys Leu Ala Asp Leu Arg Tyr Leu Ser Arg Gly Ala Ser Gly Thr Val
                                        25
                      20
          Ser Ser Ala Arg His Ala Asp Trp Arg Val Gln Val Ala Val Lys His
                              40
           35
          Leu His Ile His Thr Pro Leu Leu Asp Ser Glu Arg Lys Asp Val Leu
                                55
          Arg Glu Ala Glu Ile Leu His Lys Ala Arg Phe Ser Tyr Ile Leu Pro
                                               75
                          70
          Ile Leu Gly Ile Cys Asn Glu Pro Glu Phe Leu Gly Ile Val Thr Glu
                         85
                                           90
20
          Tyr Met Pro Asn Gly Ser Leu Asn Glu Leu Leu His Arg Lys Thr Glu
                    100
                               105
                                                110
          Tyr Pro Asp Val Ala Trp Pro Leu Arg Phe Arg Ile Leu His Glu Ile
115 120 125
          Ala Leu Gly Val Asn Tyr Leu His Asn Met Thr Pro Pro Leu Leu His
            130
                               135
                                                140
25
          His Asp Leu Lys Thr Gln Asn Ile Leu Leu Asp Asn Glu Phe His Val
                           150
                                              155
          Lys Ile Ala Asp Phe Gly Leu Ser Lys Trp Arg Met Met Ser Leu Ser
                                                    175
                        165
                                         170
          Gln Ser Arg Ser Ser Lys Ser Ala Pro Glu Gly Gly Thr Ile Ile Tyr
180 185 190
                                                       190
                     180
                                      185
30
          Met Pro Pro Glu Asn Tyr Glu Pro Gly Gln Lys Ser Arg Ala Ser Ile
                                    200
                                                    205
          Lys His Asp Ile Tyr Ser Tyr Ala Val Ile Thr Trp Glu Val Leu Ser
                                215
                                                 220
          Arg Lys Gln Pro Phe Glu Asp Val Thr Asn Pro Leu Gln Ile Met Tyr
                         230
                                             235
35
          Ser Val Ser Gln Gly His Arg Pro Val Ile Asn Glu Glu Ser Leu Pro
                        245
                                  250
          Tyr Asp Ile Pro His Arg Ala Arg Met Ile Ser Leu Ile Glu Ser Gly
                     260
                                       265
                                                        270
          Trp Ala Gln Asn Pro Asp Glu Arg Pro Ser Phe Leu Lys Cys Leu Ile
                                   280
40
          Glu Leu Glu Pro Val Leu Arg Thr Phe Glu Glu Ile Thr Phe Leu Glu
                                295
                                                  300
          Ala Val Ile Gln Leu Lys Lys Thr Lys Leu Gln Ser Val Ser Ser Ala
                          310
                                         315
          Ile His Leu Cys Asp Lys Lys Met Glu Leu Ser Leu Asn Ile Pro
                         325
                                           330
45
          Val Asn His Gly Pro Gln Glu Glu Ser Cys Gly Ser Ser Gln Leu His
                                       345
          Glu Asn Ser Gly Ser Pro Glu Thr Ser Arg Ser Leu Pro Ala Pro Gln
                                   360
                                                      365
          Asp Asn Asp Phe Leu Ser Arg Lys Ala Gln Asp Cys Tyr Phe Met Lys
                      375
                                                380
50
          Leu His His Cys Pro Gly Asn His Ser Trp Asp Ser Thr Ile Ser Gly
                         390
                                               395
          Ser Gln Arg Ala Ala Phe Cys Asp His Lys Thr Thr Pro Cys Ser Ser
                        405
                                          410
          Ala Ile Ile Asn Pro Leu Ser Thr Ala Gly Asn Ser Glu Arg Leu Gln
                     420
                                       425
55
          Pro Gly Ile Ala Gln Gln Trp Ile Gln Ser Lys Arg Glu Asp Ile Val
```

		435			440					445					
	Asn Gln 450	Met Thr	Glu Ala	Cys 455	Leu	Asn	Gln	Ser	Leu 460	Asp	Ala	Leu	Leu		
5	Ser Arg		470					475					480		
	Pro Thr	_	485				490					495			
	Gln Gly	Glu Glu 500	Phe Ala	Lys	Val	Ile 505	Val	Gln	Lys	Leu	Lys 510	Asp	Asn		
10	Lys Gln	Met Gly 515	Leu Gln	Pro	Tyr 520	Pro	Glu	Ile	Leu	Val 525	Val	Ser	Arg		
	Ser Pro 530	Ser Leu	Asn Leu	Leu 535	Gln	Asn	Lys	Ser	Met 540						
15					SEÇ	UENC	E LI	STIN	G						
	<110> Axxima Pharmaceuticals AG <120> CELLULAR KINASES INVOLVED IN CYTOMEGALOVIRUS INFECTION														
20			KINASES INHIBIT		LVED	IN (CYTO	1EGA1	LOVI	RUS :	INFE	CTION	1		
	<130> U5	0062													
	<140> <141>	s													
25	<160> 2														
	<170> Pa	tentIn V	Ver. 2.1												
30	<210> 1 <211> 26 <212> DN <213> Ho	A.	ens												
35	<220> <221> CD <222> (1		5)												
	<400> 1 atg caa Met Gln 1													48	
40	ttc ctg Phe Leu		Ala Glu		Asp	Ser	Gly			Gly		Val		96	
45	ctg tgt Leu Cys													144	
	aag ggg Lys Gly 50													192	
50	aag atg Lys Met 65													240	
55	gtc atc Val Ile													288	

5	aag Lys	ggc Gly	aac Asn	ctg Leu 100	atg Met	cac His	gtg Val	ctg Leu	aaa Lys 105	gcc Ala	gag Glu	atg Met	agt Ser	act Thr 110	ccg Pro	ctt Leu	336
	tct Ser	gta Val	aaa Lys 115	gga Gly	agg Arg	ata Ile	att Ile	ttg Leu 120	gaa Glu	atc Ile	att Ile	gaa Glu	gga Gly 125	atg Met	tgc Cys	tac Tyr	384
10	tta Leu	cat His 130	gga Gly	aaa Lys	ggc Gly	gtg Val	ata Ile 135	cac His	aag Lys	gac Asp	ctg Leu	aag Lys 140	cct Pro	gaa Glu	aat Asn	atc Ile	432
15	ctt Leu 145	gtt Val	gat Asp	aat Asn	gac Asp	ttc Phe 150	cac His	att Ile	aag Lys	atc Ile	gca Ala 155	gac Asp	ctc Leu	ggc Gly	ctt Leu	gcc Ala 160	480
		ttt Phe															528
20		gaa Glu															576
25		gcg Ala															624
	_	gat Asp 210			-		-	_	_						-		672
30		gag Glu															720
35		aaa Lys															768
		aga Arg															816
40		gct Ala		_						_	_						864
45		tta Leu 290															912
		aaa Lys															960
50	Leu	caa Gln	Leu	Asp	Cys 325	Val	Āla	Val	Pro	Ser 330	Ser	Arg	Ser	Asn	Ser 335	Ala	1008
55		gaa Glu															1056

5	cct Pro	gtg Val	gag Glu 355	gag Glu	tcc Ser	tgg Trp	ttt Phe	gct Ala 360	cct Pro	tcc Ser	ctg Leu	gag Glu	cac His 365	cca Pro	caa Gln	gaa Glu	1104
	gag Glu	aat Asn 370	gag Glu	ccc Pro	agc Ser	ctg Leu	cag Gln 375	agt Ser	aaa Lys	ctc Leu	caa Gln	gac Asp 380	gaa Glu	gcc Ala	aac As n	tac Tyr	1152
10	cat His 385	ctt Leu	tat Tyr	ggc Gly	agc Ser	cgc Arg 390	atg Met	gac Asp	agg Arg	cag Gln	acg Thr 395	aaa Lys	c ag Gln	cag Gln	ccc Pro	aga Arg 400	1200
15	cag Gln	aat Asn	gtg Val	gct Ala	tac Tyr 405	aac Asn	aga Arg	gag Glu	gag Glu	gaa Glu 410	agg Arg	aga Arg	cgc Arg	agg Arg	gtc Val 415	tcc Ser	1248
	cat His	gac Asp	cct Pro	ttt Phe 420	gca Ala	cag Gln	caa Gln	aga Arg	cct Pro 425	tac Tyr	gag Glu	aat Asn	ttt Phe	cag Gln 430	aat Asn	aca Thr	1296
20				ggc Gly													1344
25				ccc Pro													1392
				tta Leu													1440
30				gca Ala													1488
35				ctg Leu 500													1536
				acc Thr	~			_		_				_			1584
40				acc Thr													1632
45				gag Glu													1680
	aat Asn	acg Thr	aac Asn	ttc Phe	aaa Lys 565	gaa Glu	gag Glu	cca Pro	gct Ala	gct Ala 570	aag Lys	tac Tyr	caa Gln	gct Ala	atc Ile 575	ttt Phe	1728
50				act Thr 580													1776
55	aat Asn	ctg Leu	gga Gly 595	aag Lys	cac His	tgg Trp	aaa Lys	aac Asn 600	tgt Cys	gcc Ala	cgt Arg	aaa Lys	ctg Leu 605	ggc Gly	ttc Phe	aca Thr	1824

5	cag tot cag att gat gaa att gac cat gac tat gag cga gat gga ctg 1873 Gln Ser Gln Ile Asp Glu Ile Asp His Asp Tyr Glu Arg Asp Gly Leu 610 615 620	2
	aaa gaa aag gtt tac cag atg ctc caa aag tgg gtg atg agg gaa ggc 192 Lys Glu Lys Val Tyr Gln Met Leu Gln Lys Trp Val Met Arg Glu Gly 625 630 640	0
10	ata aag gga gcc acg gtg ggg aag ctg gcc cag gcg ctc cac cag tgt 1969 Ile Lys Gly Ala Thr Val Gly Lys Leu Ala Gln Ala Leu His Gln Cys 645 650 655	8
15	tcc agg atc gac ctt ctg agc agc ttg att tac gtc agc cag aac taa 2010 Ser Arg Ile Asp Leu Leu Ser Ser Leu Ile Tyr Val Ser Gln Asn 660 665 670	6
	ceetggatgg getaeggeag etgaagtgga egeeteaett ageggataae eecagaaagt 2070	6
	tggctgcctc agagcattca gaattctgtc ctcactgata ggggttctgt gtctgcagaa 213	6
20	atttngtttc ctgtacttca tagctggaga atggggaaag aaatctgcag caaaggggtc 2196	6
	tcactctgtt gecaggetgg tetcaaaett etggaetcaa gtgateetee egeeteggee 2256	6
	ttccaaagtg ctgggatatc aggcactgag ccactgcgcc cagtcaacaa tccgntctga 2316	5
25	ggaaagegta agcaggaaga eetettaatg geatageace aataaaaaaa tgacteetag 2376	6
	ttgtgtttgg aaagggagag aagagatgte tgaggaaggt catgttettt cagettatgg 2430	6
	catttcctag agtttngttg aagcaagaag aaaaactcag agaatataaa atcaactttn 2496	5
30	aaaattgtgt gctctcttct tcacgtaggc tcctgttaaa aacaaagtgc agtcagattc 2556	5
	taagccctgt tcagagactt cgcggatcac agctgcagct caccgccaca tcacaggatc 2616	5
	C - 2617	7
35	<210> 2 <211> 671 <212> PRT <213> Homo sapiens	
	<400> 2 Met Gln Pro Asp Met Ser Leu Asn Val Ile Lys Met Lys Ser Ser Asp	
40	1 5 10 15 Phe Leu Glu Ser Ala Glu Leu Asp Ser Gly Gly Phe Gly Lys Val Ser	
	20 25 30	
	Leu Cys Phe His Arg Thr Gln Gly Leu Met Ile Met Lys Thr Val Tyr 35 40 45 Lys Gly Bro Agn Cyg Ile Gly Hig Arg Gly Ale Lys Gly Bl	
45	Lys Gly Pro Asn Cys Ile Glu His Asn Glu Ala Leu Leu Glu Glu Ala 50 55 60 Lys Mot Mot Asn Arg Leu Arg His San Arg Val Val Val Val	
	Lys Met Met Asn Arg Leu Arg His Ser Arg Val Val Lys Leu Leu Gly 65 70 75 80	
	Val Ile Ile Glu Glu Gly Lys Tyr Ser Leu Val Met Glu Tyr Met Glu 85 90 95	
50	Lys Gly Asn Leu Met His Val Leu Lys Ala Glu Met Ser Thr Pro Leu 100 105 110 Ser Val Lya Gly Ang Ila Ila Lau Gly Ila	
	Ser Val Lys Gly Arg Ile Ile Leu Glu Ile Ile Glu Gly Met Cys Tyr 115 120 125	
	Leu His Gly Lys Gly Val Ile His Lys Asp Leu Lys Pro Glu Asn Ile 130 135 140	
55	Leu Val Asp Asn Asp Phe His Ile Lys Ile Ala Asp Leu Gly Leu Ala 145 150 155 160	

			_		_		_		3	3	a 1	a 1	uia	Nan	C1	Lou
		Phe	-		165					170					175	
5	Arg	Glu	Val	Asp 180	Gly	Thr	Ala	Lys	Lys 185	Asn	Gly	Gly	Thr	Leu 190	Tyr	Tyr
	Met	Ala	Pro 195	Glu	His	Leu	Asn	Asp 200	Val	Asn	Ala	Lys	Pro 205	Thr	Glu	Lys
	Ser	Asp 210	Val	Tyr	Ser	Phe	Ala 215	Val	Val	Leu	Trp	Ala 220	Ile	Phe	Ala	Asn
10	Lys 225	Glu	Pro	Tyr	Glu	Asn 230	Ala	Ile	Cys	Glu	Gln 235	Gln	Leu	Ile	Met	Cys 240
		Lys	Ser	Gly	Asn 245	Arg	Pro	Asp	Val	Asp 250	Asp	Ile	Thr	Glu	Tyr 255	Cys
	Pro	Arg	Glu	Ile 260	Ile	Ser	Leu	Met	Lys 265	Leu	Cys	Trp	Glu	Ala 270	Asn	Pro
15	Glu	Ala	Arg 275	Pro	Thr	Phe	Pro	Gly 280	Ile	Glu	Glu	Lys	Phe 285	Arg	Pro	Phe
	-	Leu 290					295					300				
	305	Lys		_		310					315					320
20		Gln		-	325					330					335	
		Glu		340	-				345			_		350		_
		Val	355			_		360					365			
25		Asn 370					375		-			380				_
	385	Leu	-	_		390		-	_		395	_				400
		Asn			405					410	_	_		_	415	
30		Asp		420				_	425	_				430		
		Gly His	435	-			-	440					445	_		
		450 Asn				_	455					460			_	
35	465		_		_	470			_		475		_			480
		Gly			485				_	490					495	
		Pro Thr		500					505					510		_
40		Lys	515					520					525	-		
		530 Tyr	_				535			_		540		_		_
45	545	Thr				550	_				555			_		560
45		Asn			565					570					575	
		Leu		580					585					590		
50		Ser	595					600				-	605	_		
		610 Glu					615					620				
	625 Ile	Lys	Gly	Ala		630 Val	Gly	Lys	Leu	Ala	635 Gln	Ala	Leu	His	Gln	640 Cys
55	ser	Arg	Ile	Asp	645 Leu	Leu	Ser	Ser	Leu	650 Ile	Tyr	Val	Ser	Gln	655 Asn	
				660					665					670		

SEQUENCE LISTING

5	<110> Axxima Pharmaceuticals AG
	<120> CELLULAR KINASES INVOLVED IN CYTOMEGALOVIRUS INFECTION AND THEIR INHIBITION
10	<130> Y10256
	<140> <141>
	<160> 2
15	<170> PatentIn Ver. 2.1
20	<210> 1 <211> 4596 <212> DNA <213> Homo sapiens
	<220> <221> CDS <222> (233)(3076)
25	<400> 1 aagcggggga ctgtgccgtg tggaacgtgt agctgttgag aggtggactc tgttaccatt 60
	gaggatgttt ggaggatgag tatgtgtggc agaggcacac ataaacaggc agagaccctt 120
	tgeccetgee tttetcccce aacceaagge tgacctgtgt tetcccaggt ctgggattet 180
30	aagtgacetg etetgtgttt ggtetetete aggatgagea caageetggg ag atg gea 238 Met Ala 1
35	gtg atg gaa atg gcc tgc cca ggt gcc cct ggc tca gca gtg ggg cag Val Met Glu Met Ala Cys Pro Gly Ala Pro Gly Ser Ala Val Gly Gln 5 10 15
	cag aag gaa ctc ccc aag cca aag gag aag acg ccg cca ctg ggg aag 334 Gln Lys Glu Leu Pro Lys Pro Lys Glu Lys Thr Pro Pro Leu Gly Lys 20 25 30
40	aaa cag agc tcc gtc tac aag ctt gag gcc gtg gag aag agc cct gtg Lys Gln Ser Ser Val Tyr Lys Leu Glu Ala Val Glu Lys Ser Pro Val 35 40 45 50
45	ttc tgc gga aag tgg gag atc ctg aat gac gtg att acc aag ggc aca Phe Cys Gly Lys Trp Glu Ile Leu Asn Asp Val Ile Thr Lys Gly Thr 55 60 65
	gcc aag gaa ggc tcc gag gca ggg cca gct gcc atc tct atc atc gcc 478 Ala Lys Glu Gly Ser Glu Ala Gly Pro Ala Ala Ile Ser Ile Ile Ala 70 75 80
50	cag gct gag tgt gag aat agc caa gag ttc agc ccc acc ttt tca gaa 526 Gln Ala Glu Cys Glu Asn Ser Gln Glu Phe Ser Pro Thr Phe Ser Glu 85 90 95
55	cgc att ttc atc gct ggg tcc aaa cag tac agc cag tcc gag agt ctt 574 Arg Ile Phe Ile Ala Gly Ser Lys Gln Tyr Ser Gln Ser Glu Ser Leu 100 105 110

5	gat Asp 115	cag Gln	atc Ile	ccc Pro	aac Asn	aat Asn 120	gtg Val	gcc Ala	cat His	gct Ala	aca Thr 125	gag Glu	ggc Gly	aaa Lys	atg Met	gcc Ala 130	622
	cgt Arg	gtg Val	tgt Cys	tgg Trp	aag Lys 135	gga Gly	aag Lys	cgt Arg	cgc Arg	agc Ser 140	aaa Lys	gcc Ala	cgg Arg	aag Lys	aaa Lys 145	cgg Arg	670
10	aag Lys	aag Lys	aag Lys	agc Ser 150	tca Ser	aag Lys	tcc Ser	ctg Leu	gct Ala 155	cat His	gca Ala	gga Gly	gtg Val	gcc Ala 160	ttg Leu	gcc Ala	718
15	aaa Lys	ccc Pro	ctc Leu 165	ccc Pro	agg Arg	acc Thr	cct Pro	gag Glu 170	cag Gln	gag Glu	agc Ser	tgc Cys	acc Thr 175	atc Ile	cca Pro	gtg Val	766
	cag Gln	gag Glu 180	gat Asp	gag Glu	tct Ser	cca Pro	ctc Leu 185	ggc Gly	gcc Ala	cca Pro	tat Tyr	gtt Val 190	aga Arg	aac Asn	acc Thr	ccg Pro	814
20		ttc Phe															862
25		cag Gln															910
		aaa Lys															958
30		cac His															1006
35		tat Tyr 260	_	_	-										_		1054
40		aaa Lys															1102
40		agc Ser															1150
45		gac As p															1198
50		tct Ser															1246
50		gct Ala 340															1294
55	agc Ser 355	ctg Leu	gcc Ala	aag Lys	acc Thr	tgg Trp 360	gca Ala	gca Ala	cgg Arg	ggc Gly	tcc Ser 365	aga Arg	tcc Ser	cgg Arg	gag Glu	ccc Pro 370	1342

5	agc Ser	ccc Pro	aaa Lys	act Thr	gag Glu 375	gac Asp	aac Asn	gag Glu	ggt Gly	gtc Val 380	ctg Leu	ctc Leu	act Thr	gag Glu	aaa Lys 385	ctc Leu	1390
	aag Lys	cca Pro	gtg Val	gat Asp 390	tat Tyr	gag Glu	tac Tyr	cga Arg	gaa Glu 395	gaa Glu	gtc Val	cac His	tgg Trp	gcc Ala 400	acg Thr	cac His	1438
10	cag Gln	ctc Leu	cgc Arg 405	ctg Leu	ggc Gly	aga Arg	ggc Gly	tcc Ser 410	ttc Phe	gga Gly	gag Glu	gtg Val	cac His 415	agg Arg	atg Met	gag Glu	1486
15	gac Asp	aag Lys 420	cag Gln	act Thr	ggc Gly	ttc Phe	cag Gln 425	tgc Cys	gct Ala	gtc Val	aaa Lys	aag Lys 430	gtg Val	cgg Arg	ctg Leu	gaa Glu	1534
	gta Val 435	ttt Phe	cgg Arg	gca Ala	gag Glu	gag Glu 440	ctg Leu	atg Met	gca Ala	tgt Cys	gca Ala 445	gga Gly	ttg Leu	acc Thr	tca Ser	ccc Pro 450	1582
20				cct Pro													1630
25				gag Glu 470													1678
30				tgt Cys													1726
30				ggt Gly													1774
35	_	~		gct Ala	_				_		-	-		_			1822
40				gac Asp													1870
70		_		ttg Leu 550				-									1918
45				gag Glu													1966
50				agc Ser													2014
				cag Gln													2062
55	gag Glu	cct Pro	ccg Pro	cct Pro	gtg Val 615	agg Arg	gag Glu	atc Ile	cca Pro	ccc Pro 620	tcc Ser	tgc Cys	gcc Ala	cct Pro	ctc Leu 625	aca Thr	2110

5	gcc Ala	cag Gln	gcc Ala	atc Ile 630	caa Gln	gag Glu	Gly aaa	ctg Leu	agg Arg 635	aaa Lys	gag Glu	ccc Pro	atc Ile	cac His 640	cgc Arg	gtg Val	2158
	tct Ser	gca Ala	gcg Ala 645	gag Glu	ctg Leu	gga Gly	Gly aaa	aag Lys 650	gtg Val	aac Asn	cgg Arg	gca Ala	cta Leu 655	cag Gln	caa Gln	gtg Val	2206
10	gga Gly	ggt Gly 660	ctg Leu	aag Lys	agc Ser	cct Pro	tgg Trp 665	agg Arg	gga Gly	gaa Glu	tat Tyr	aaa Lys 670	gaa Glu	cca Pro	aga Arg	cat His	2254
15	cca Pro 675	ccg Pro	cca Pro	aat Asn	caa Gln	gcc Ala 680	aat Asn	tac Tyr	cac His	cag Gln	acc Thr 685	ctc Leu	cat His	gcc Ala	cag Gln	ccg Pro 690	2302
				tcg Ser													2350
20				gcc Ala 710													2398
25				aag Lys				_		_	-	_					2446
20				ccc Pro													2494
30				tca Ser													2542
35	_	_	_	gaa Glu		_					_	_		_			2590
40				gag Glu 790													2638
40			_	tcg Ser	_	~	~	-	_				_	_			2686
45				gac Asp													2734
50				cga Arg													2782
				acc Thr													2830
55	tct Ser	ctt Leu	aat Asn	ggt Gly 870	gaa Glu	cac His	ctg Leu	cac His	atc Ile 875	cgg Arg	gag Glu	ttc Phe	cac His	cgg Arg 880	gtc Val	aaa Lys	2878

5	gtg gga gac atc gcc act ggc atc agc agc cag atc cca gct gca gcc 2920 Val Gly Asp Ile Ala Thr Gly Ile Ser Ser Gln Ile Pro Ala Ala Ala 885 890 895	5
	ttc agc ttg gtc acc aaa gac ggg cag cct gtt cgc tac gac atg gag 297. Phe Ser Leu Val Thr Lys Asp Gly Gln Pro Val Arg Tyr Asp Met Glu 900 905 910	4
10	gtg cca gac tcg ggc atc gac ctg cag tgc aca ctg gcc cct gat ggc 302. Val Pro Asp Ser Gly Ile Asp Leu Gln Cys Thr Leu Ala Pro Asp Gly 915 920 925 930	2
15	agc ttc gcc tgg agc tgg agg gtc aag cat ggc cag ctg gag aac agg 307 Ser Phe Ala Trp Ser Trp Arg Val Lys His Gly Gln Leu Glu Asn Arg 935 940 945	0
	ccc taa ccctgccctc caccgccggc tccacactgc cggaaagcag ccttcctgct 312 Pro	6
20	eggtgcacga tgetgeeetg aaaacacagg etcageegtt eecaggggat tgecageeec 318	6
	ccggctcaca gtgggaacca gggcctcgca gcagcaaggt gggggcaagc agaatgcctc 324	6
	ccaggattte acacetgage ectgeeccae ectgetgaaa aaacateege caegtgaaga 330	6
25	gacagaagga ggatggcagg agttacctgg ggaaacaaaa cagggatctt tttctgcccc 336	6
	tgctccagtc gagttggcct gacccgcttg gatcagtgac catttgttgg cagacagggg 342	б
	agagcagett ceageetggg teagaagggg tgggegagee etteggeece teaceeteea 348	б
30	ggctgctgtg agagtgtcaa gtgtgtaagg gcccaaactc aggttcagtg cagaaccagg 354	6
	tcagcaggta tgcccgcccg taggttaagg gggccctcta aaccccttgc ctggcctcac 3600	
	ctggccagct cacccctttt gggtgtaggg gaaaagaatg cctgaccctg ggaaggctcc 366	
35	ctggtagaat acaccacact tttcaggttg ttgcaacaca ggtcctgagt tgacctctgg 372	5
	ttcagccaag gaccaaagaa ggtgtgtaag tgaagtggtt ctcagtcccc agacatgtgc 378	5
	ccctttgctg ctggctacca ctcttcccca gagcagcagg ccccgagccc cttcaggccc 384	5
40	agcactgccc cagactcgct ggcactcagt teceteatet gtaaaggtga agggtgatge 3900	5
	aggatatgcc tgacaggaac agtctgtgga tggacatgat cagtgctaag gaaagcagca 396	5
	gagagagacg teeggegeee cageeeeact ateagtgtee agegtgetgg tteeceagag 4020	5
45	cacageteag cateacastg acasteases tgesetgess stggssagag ggtastgssg 408	6
	acggeaettt geaetetgat gaeeteaaag eaettteatg getgeeetet ggeagggeag	6
	ggcagggcag tgacactgta ggagcatagc aagccaggag atggggtgaa gggacacagt 4200	5
50	cttgagctgt ccacatgcat gtgactcctc aaacctcttc cagatttctc taagaatagc 426	5
	accecettee ceattgeece agettageet etteteecag gggagetaet caggaeteae 432	6
	gtagcattaa atcagctgtg aatcgtcagg gggtgtctgc tagcctcaac ctcctggggc 4380	
55	aggggacgee gagaeteegt gggagaaget catteecaca tettgecaag acageetttg 4440	5

tocagetgte cacattgagt cagactgete ceggggagag ageceeggee cecageacat 4506 aaagaactgc agccttggta ctgcagagtc tgggttgtag agaactcttt gtaagcaata 4566 aaqtttqqqq tqatgacaaa tgttaaaaaa <210> 2 <211> 947 <212> PRT <213> Homo sapiens <400>2Met Ala Val Met Glu Met Ala Cys Pro Gly Ala Pro Gly Ser Ala Val Gly Gln Gln Lys Glu Leu Pro Lys Pro Lys Glu Lys Thr Pro Pro Leu 2.0 Gly Lys Lys Gln Ser Ser Val Tyr Lys Leu Glu Ala Val Glu Lys Ser Pro Val Phe Cys Gly Lys Trp Glu Ile Leu Asn Asp Val Ile Thr Lys 5.5 Gly Thr Ala Lys Glu Gly Ser Glu Ala Gly Pro Ala Ala Ile Ser Ile Ile Ala Gln Ala Glu Cys Glu Asn Ser Gln Glu Phe Ser Pro Thr Phe Ser Glu Arg Ile Phe Ile Ala Gly Ser Lys Gln Tyr Ser Gln Ser Glu Ser Leu Asp Gln Ile Pro Asn Asn Val Ala His Ala Thr Glu Gly Lys Met Ala Arg Val Cys Trp Lys Gly Lys Arg Arg Ser Lys Ala Arg Lys Lys Arg Lys Lys Ser Ser Lys Ser Leu Ala His Ala Gly Val Ala Leu Ala Lys Pro Leu Pro Arg Thr Pro Glu Gln Glu Ser Cys Thr Ile Pro Val Gln Glu Asp Glu Ser Pro Leu Gly Ala Pro Tyr Val Arg Asn Thr Pro Gln Phe Thr Lys Pro Leu Lys Glu Pro Gly Leu Gly Gln Leu Cys Phe Lys Gln Leu Gly Glu Gly Leu Arg Pro Ala Leu Pro Arg Ser Glu Leu His Lys Leu Ile Ser Pro Leu Gln Cys Leu Asn His Val Trp Lys Leu His His Pro Gln Asp Gly Gly Pro Leu Pro Leu Pro Thr His Pro Phe Pro Tyr Ser Arg Leu Pro His Pro Phe Pro Phe His Pro Leu Gln Pro Trp Lys Pro His Pro Leu Glu Ser Phe Leu Gly Lys Leu Ala Cys Val Asp Ser Gln Lys Pro Leu Pro Asp Pro His Leu Ser Lys Leu Ala Cys Val Asp Ser Pro Lys Pro Leu Pro Gly Pro His Leu Glu Pro Ser Cys Leu Ser Arg Gly Ala His Glu Lys Phe Ser Val Glu Glu Tyr Leu Val His Ala Leu Gln Gly Ser Val Ser Ser Ser Gln Ala His Ser Leu Thr Ser Leu Ala Lys Thr Trp Ala Ala Arg Gly Ser Arg Ser Arg Glu Pro Ser Pro Lys Thr Glu Asp Asn Glu Gly Val Leu Leu Thr Glu Lys Leu Lys Pro Val Asp Tyr Glu Tyr Arg Glu Glu Val His Trp Ala Thr His Gln Leu Arg Leu Gly Arg Gly Ser Phe Gly Glu Val His Arg

					405		_	_,		410	- 1		_	-	415	
	Met	Glu	Asp		Gln	Thr	Gly	Phe		Cys	Ala	Val	Lуs	Lуs 430	Val	Arg
5	Lou	Glu	17a l	420 Dhe	Nrα	λla	Glu	Glu	425 Leu	Met	Δla	Cvs	Ala		Leu	Thr
	Leu	GIU	435	FIIC	Arg	AIG	Olu	440	цец		111.0	CIO	445			
	Ser	Pro		Ile	Val	Pro	Leu	Tyr	Gly	Ala	Val	Arg	${\tt Glu}$	Gly	Pro	Trp
		450					455	_	~1	~ 7 .	a 1	460	T	a1	a1	T
		Asn	Ile	Phe	Met	G1u 470	Leu	Leu	GIU	GIY	475	ser	ьeu	GIY	GIN	480
10	465 Val	Lys	Glu	Gln	Glv		Leu	Pro	Glu	Asp		Ala	Leu	Tyr	Tyr	
		-			485					490					495	
	Gly	Gln	Ala		Glu	Gly	Leu	Glu		Leu	His	Ser	Arg		Ile	Leu
	ніе	Gly	λen	500 Val	Lve	Δla	Agn	Asn	505 Val	Len	Leu	Ser	Ser	510 Asp	Glv	Ser
15	птэ	Gry	515	val	БуБ	ma	7100	520		200	200	201	525	<u>-</u>	1	
	His	Ala	Ala	Leu	Cys	Asp	Phe	Gly	His	Ala	Val	Cys	Leu	Gln	Pro	Asp
	G3	530	a 1	T	0	T 011	535	mb w	G1	n an	Tree.	540	Dro	C1.,	Thr	Clu
	545	Leu	GIY	гув	ser	550	ьеи	THE	GTÀ	Asp	555	TIE	PLO	GIY	TIIL	560
		His	Met	Ala	Pro		Val	Val	Leu	Gly		Ser	Cys	Asp	Ala	-
20					565					570					575	
	Val	Asp	Val	Trp 580	Ser	Ser	Cys	Cys	Met 585	Met	Leu	His	Met	Leu 590	Asn	GIÀ
	Cvs	His	Pro		Thr	Gln	Phe	Phe		Gly	Pro	Leu	Cys		Lys	Ile
	-1-		595					600					605		-	
	Ala	Ser	Glu	Pro	Pro	Pro		Arg	Glu	Ile	Pro		Ser	Суѕ	Ala	Pro
25	T 011	610 Thr	. ד ת	Cln	~ ו ת	Tla	615 Cln	0111	C111	LOU	A ra	620	G1,,	Dro	Tla	ui e
	625	1111	AIa	Gill	AIG	630	GIII	Giu	GLY	пец	635	шyз	Olu	110	110	640
	Arg	Val	Ser	Ala	Ala	Glu	Leu	Gly	Gly	Lys	Val	Asn	Arg	Ala	Leu	Gln
	~1	T V - 3	~1	a1	645	T	0	D	m	650	a1	a1	m	* =	655	D
30	GIN	Val	GIY	660	Leu	гÀг	ser	Pro	665	Arg	GIY	GIU	туr	ьуs 670	GIU	Pro
	Arg	His	Pro		Pro	Asn	Gln	Ala		Tyr	His	Gln	Thr		His	Ala
			675				_	680		_			685	_		
	GIn	Pro 690	Arg	GIu	Leu	Ser	Pro 695	Arg	Ala	Pro	GIA	700	Arg	Pro	Ala	GLu
	Glu	Thr	Thr	Gly	Arg	Ala		Lys	Leu	Gln	Pro		Leu	Pro	Pro	Glu
35	705			-	•	710		-			715					720
	Pro	Pro	Glu	Pro		Lys	Ser	Pro	Pro		Thr	Leu	Ser	Lys		Glu
	Ser	Gly	Met	Trp	725 Glu	Pro	Leu	Pro	Leu	730 Ser	Ser	Leu	Glu	Pro	735 Ala	Pro
		1		740					745					750	-12-0	
	Ala	Arg		Pro	Ser	Ser	Pro		Arg	Lys	Ala	Thr		Pro	Glu	Gln
40	Glu	Leu	755	Gln	T.eu	Glu	Tle	760	T.e.11	Dhe	T.e.11	Δen	765 Ser	Len	Sar	Gln
	GIG	770	GIII	0111	шси	Olu	775	Gra	Lea	FIIC	пса	780	Der	пец	Der	GIII
	Pro	Phe	Ser	Leu	${\tt Glu}$	${\tt Glu}$	Gln	${\tt Glu}$	Gln	Ile	Leu	Ser	Cys	Leu	Ser	Ile
	785	602	T 011	Cor	Low	790	λαν) an	Cox	~1. ,	795	7.00	Dwo	0.02	*	800
45	Asp	Ser	пеп	ser	805	ser	Asp	Asp	Ser	810	ьуѕ	ASII	Pro	ser	815	Ald
	Ser	Gln	Ser	Ser		Asp	Thr	Leu	Ser		Gly	Val	His	Ser		Ser
		~ 3		820		_	_	_	825		_			830		_
	ser	Gln	A1a 835	Glu	Ala	Arg	Ser	Ser 840	Ser	Trp	Asn	Met	Va⊥ 845	Leu	Ala	Arg
	Gly	Arg		Thr	Asp	Thr	Pro		Tyr	Phe	Asn	Gly		Lys	Val	Gln
50		850			_		855		_			860		-		
		Gln	Ser	Leu	Asn		Glu	His	Leu	His		Arg	Glu	Phe	His	_
	865 Val	Lys	Val	Glv	Asn	870 Ile	Ala	Thr	Glv	Πle	875 Ser	Ser	Gln	Ile	Pro	880 Ala
					885					890					895	
	Ala	Ala	Phe		Leu	Val	Thr	Lys		Gly	Gln	Pro	Val		Tyr	Asp
55	Met	Glu	Val	900 Pro	Asn	Ser	Glv	Tle	905 Agn	Lev	Gln	Cve	Thr	910	λla	Dro
		- Lu		0	nsp		Gry	116	тар	ьeu	3111	CYB	TILL	neu	Ala	F10

5	915 920 925 Asp Gly Ser Phe Ala Trp Ser Trp Arg Val Lys His Gly Gln Leu Glu 930 935 940 Asn Arg Pro 945
10	SEQUENCE LISTING
	<110> Axxima Pharmaceuticals AG
15	<pre><120> CELLULAR KINASES INVOLVED IN CYTOMEGALOVIRUS INFECTION AND THEIR INHIBITION <130> NM002756</pre>
	<140>
	<141>
20	<160> 2
	<170> PatentIn Ver. 2.1
25	<210> 1 <211> 2030 <212> DNA <213> Homo sapiens
30	<220> <221> CDS <222> (338)(1294)
	<400> 1 tggctggcaa tggccttgct gacctcgagc cgggcccacg tggggacctt tggagcacag 60
	cctacgatcc tggtgcaagg ccggtggatg cagaggccag tccatatacc acccaggcct 120
35	gcgaggagcg tggtccccac ccatccagcc catatgtgca agtgccettg acagagagc 180
	tggtcatatc catggtgacc atttatgggc cacaacaggt ccccatetgc gcagtgaacc 240
	ctgtgctgag caccttgcag acgtgatctt gcttcgtcct gcagcactgt gcggggcagg 300
40	aaaatccaag aggaagaagg atctacggat atcctgc atg tcc aag cca ccc gca 355 Met Ser Lys Pro Pro Ala 1 5
45	ccc aac ccc aca ccc ccc cgg aac ctg gac tcc cgg acc ttc atc acc Pro Asn Pro Thr Pro Pro Arg Asn Leu Asp Ser Arg Thr Phe Ile Thr 10 15 20
	att gga gac aga aac ttt gag gtg gag gct gat gac ttg gtg acc atc 11e Gly Asp Arg Asn Phe Glu Val Glu Ala Asp Asp Leu Val Thr Ile 25 30 35
50	tca gaa ctg ggc cgt gga gcc tat ggg gtg gta gag aag gtg cgg cac Ser Glu Leu Gly Arg Gly Ala Tyr Gly Val Val Glu Lys Val Arg His 40 45 50
55	gcc cag agc ggc acc atc atg gcc gtg aag cgg atc cgg gcc acc gtg Ala Gln Ser Gly Thr Ile Met Ala Val Lys Arg Ile Arg Ala Thr Val 60 65 70

5	aac Asn	tca Ser	cag Gln	gag Glu	cag Gln 75	aag Lys	cgg Arg	ctg Leu	ctc Leu	atg Met 80	gac Asp	ctg Leu	gac Asp	atc Ile	aac Asn 85	atg Met	595
	cgc Arg	acg Thr	gtc Val	gac Asp 90	tgt Cys	ttc Phe	tac Tyr	act Thr	gtc Val 95	acc Thr	ttc Phe	tac Tyr	Gly 999	gca Ala 100	cta Leu	ttc Phe	643
10	aga Arg	gag Glu	gga Gly 105	gac Asp	gtg Val	tgg Trp	atc Ile	tgc Cys 110	atg Met	gag Glu	ctc Leu	atg Met	gac Asp 115	aca Thr	tcc Ser	ttg Leu	691
15	gac Asp	aag Lys 120	ttc Phe	tac Tyr	cgg Arg	aag Lys	gtg Val 125	ctg L eu	gat Asp	aaa Lys	aac Asn	atg Met 130	aca Thr	att Ile	cca Pro	gag Glu	739
	gac Asp 135	atc Ile	ctt Leu	Gly aaa	gag Glu	att Ile 140	gct Ala	gtg Val	tct Ser	atc Ile	gtg Val 145	cgg Arg	gcc Ala	ctg Leu	gag Glu	cat His 150	787
20		cac His															835
25	_	ctt Leu			_					-	_	_		_		_	883
		ggc Gly															931
30		ccc Pro 200															979
35		tac Tyr															1027
		atg Met	_		_										_		1075
40	_	cag Gln	_	_	_					_			-			_	1123
45		cgt Arg															1171
	-	aac Asn 280		-		_	-	-		_		_	-				1219
50		ttc Phe															1267
55		aag Lys							tag	gggo	etggg	jcc t	cgga	ccc	ca		1314

```
ctccggccct ccagagcccc acagccccat ctgcgggggc agtgctcacc cacaccataa 1374
           getactgeca tectggecca gggcatetgg gaggaacega gggggetget eccaeetgge 1434
5
           tetgtggega gecatttgte ecaagtgeea aagaageaga ecattgggge teecagecag 1494
           geoettgteg geoecaecag tgeeteteec tgetgeteet aggaceegte tecagetget 1554
           gagatectgg actgaggggg cetggatgee eeetgtggat getgetgeee etgeacagea 1614
10
           ggctgccagt gcctgggtgg atgggccacc gccttgccca gcctggatgc catccaagtt 1674
           gtatattttt ttaatctctc gactgaatgg actttgcaca ctttggccca gggtggccac 1734
           acctctatcc cggctttggt gcggggtaca caagagggga tgagttgtgt gaatacccca 1794
15
           agacteccat gagggagatg ccatgageeg eccaaggeet teecetggea etggcaaaca 1854
           qqqcctctgc ggagcacact ggctcaccca gtcctgcccg ccaccgttat cggtgtcatt 1914
           cacctttcgt gtttttttta atttatcctc tgttgatttt ttcttttgct ttatgggttt 1974
20
           ggettgtttt tettgeatgg tttggagetg ategettete ecceaecece tagggg
           <210> 2
25
           <211> 318
           <212> PRT
           <213> Homo sapiens
           <400> 2
           Met Ser Lys Pro Pro Ala Pro Asn Pro Thr Pro Pro Arg Asn Leu Asp
30
                                                 10
           Ser Arg Thr Phe Ile Thr Ile Gly Asp Arg Asn Phe Glu Val Glu Ala
                        20
                                            25
           Asp Asp Leu Val Thr Ile Ser Glu Leu Gly Arg Gly Ala Tyr Gly Val
                    35
                                        40
                                                             45
           Val Glu Lys Val Arg His Ala Gln Ser Gly Thr Ile Met Ala Val Lys
35
                50
                                    55
           Arg Ile Arg Ala Thr Val Asn Ser Gln Glu Gln Lys Arg Leu Leu Met
                                70
            65
                                                     75
           Asp Leu Asp Ile Asn Met Arg Thr Val Asp Cys Phe Tyr Thr Val Thr
                            85
                                                 90
                                                                     95
           Phe Tyr Gly Ala Leu Phe Arg Glu Gly Asp Val Trp Ile Cys Met Glu
40
                       100
                                           105
                                                                110
           Leu Met Asp Thr Ser Leu Asp Lys Phe Tyr Arg Lys Val Leu Asp Lys
                   115
                                        120
                                                            125
           Asn Met Thr Ile Pro Glu Asp Ile Leu Gly Glu Ile Ala Val Ser Ile
               130
                                   135
                                                        140
           Val Arg Ala Leu Glu His Leu His Ser Lys Leu Ser Val Ile His Arg
45
                               150
                                                    155
           Asp Val Lys Pro Ser Asn Val Leu Ile Asn Lys Glu Gly His Val Lys
                           165
                                               170
           Met Cys Asp Phe Gly Ile Ser Gly Tyr Leu Val Asp Ser Val Ala Lys
                       180
                                            185
           Thr Met Asp Ala Gly Cys Lys Pro Tyr Met Ala Pro Glu Arg Ile Asn
50
                   195
                                        200
           Pro Glu Leu Asn Gln Lys Gly Tyr Asn Val Lys Ser Asp Val Trp Ser
               210
                                   215
                                                        220
           Leu Gly Ile Thr Met Ile Glu Met Ala Ile Leu Arg Phe Pro Tyr Glu
           225
                               230
                                                    235
           Ser Trp Gly Thr Pro Phe Gln Gln Leu Lys Gln Val Val Glu Pro
55
                           245
                                                250
           Ser Pro Gln Leu Pro Ala Asp Arg Phe Ser Pro Glu Phe Val Asp Phe
```

	260 265 270 Thr Ala Gln Cys Leu Arg Lys Asn Pro Ala Glu Arg Met Ser Tyr Leu
5	275 280 285 Glu Leu Met Glu His Pro Phe Phe Thr Leu His Lys Thr Lys Lys Thr
	290 295 300 Asp Ile Ala Ala Phe Val Lys Lys Ile Leu Gly Glu Asp Ser 305 310 315
10	SEQUENCE LISTING
	<110> Axxima Pharmaceuticals AG
15	<120> CELLULAR KINASES INVOLVED IN CYTOMEGALOVIRUS INFECTION AND THEIR INHIBITION
	<130> U 88666
	<140> <141>
20	<160> 2
	<170> PatentIn Ver. 2.1
25	<210> 1 <211> 3745 <212> DNA <213> Homo sapiens
30	<220> <221> CDS <222> (188)(2248)
	<400> 1 gaatteggea egaggeeatt gaateeeagt eetaaeagaa gtaetgegaa tettgtggee 60
	tcattetgaa caaaagggat tagagaagaa aaatetettg atataagget tgaaagcaag 120
35	ggcaggcaat cttggttgtg aatattttct gatttttcca gaaatcaagc agaagattga 180
	gctgctg atg tca gtt aac tct gag aag tcg tcc tct tca gaa agg ccg Met Ser Val Asn Ser Glu Lys Ser Ser Ser Glu Arg Pro 1 5 10
40	gag cct caa cag aaa gct cct tta gtt cct cct cct cca ccg cca cca 277 Glu Pro Gln Gln Lys Ala Pro Leu Val Pro Pro Pro Pro Pro Pro Pro 15 20 25 30
45	cca cca cca ccg cca cct ttg cca gac ccc aca ccc ccg gag cca gag Pro Pro Pro Pro Pro Leu Pro Asp Pro Thr Pro Pro Glu Pro Glu 35 40 45
	gag gag atc ctg gga tca gat gat gag gag caa gag gac cct gcg gac 373 Glu Glu Ile Leu Gly Ser Asp Asp Glu Glu Glu Glu Asp Pro Ala Asp 50 55 60
50	tac tgc aaa ggt gga tat cat cca gtg aaa att gga gac ctc ttc aat 421 Tyr Cys Lys Gly Gly Tyr His Pro Val Lys Ile Gly Asp Leu Phe Asn 65 70 75
55	ggc cgg tat cat gtt att aga aag ctt gga tgg ggg cac ttc tct act 469 Gly Arg Tyr His Val Ile Arg Lys Leu Gly Trp Gly His Phe Ser Thr 80 90

5	gtc Val 95	tgg Trp	ctg Leu	tgc Cys	tgg Trp	gat Asp 100	atg Met	cag Gln	ggg Gly	aaa Lys	aga Arg 105	ttt Phe	gtt Val	gca Ala	atg Met	aaa Lys 110	517
	gtt Val	gta Val	aaa Lys	agt Ser	gcc Ala 115	cag Gln	cat His	tat Tyr	acg Thr	gag Glu 120	aca Thr	gcc Ala	ttg Leu	gat Asp	gaa Glu 125	ata Ile	5 65
10	aaa Lys	ttg Leu	ctc Leu	aaa Lys 130	tgt Cys	gtt Val	cga Arg	gaa Glu	agt Ser 135	gat Asp	ccc Pro	agt Ser	gac Asp	cca Pro 140	aac Asn	aaa Lys	613
15	gac Asp	atg Met	gtg Val 145	gtc Val	cag Gln	ctc Leu	att Ile	gac Asp 150	gac Asp	ttc Phe	aag Lys	att Ile	tca Ser 155	ggc Gly	atg Met	aat Asn	661
20	Gly ggg	ata Ile 160	cat His	gtc Val	tgc Cys	atg Met	gtc Val 165	ttc Phe	gaa Glu	gta Val	ctt L e u	ggc Gly 170	cac His	cat His	ctc Leu	ctc Leu	709
20	aag Lys 175	tgg Trp	atc Ile	atc Ile	aaa Lys	tcc Ser 180	aac Asn	tat Tyr	caa Gln	ggc Gly	ctc Leu 185	cca Pro	gta Val	cgt Arg	tgt Cys	gtg Val 190	757
25				att Ile													805
30				atc Ile 210													853
				gat Asp													901
35				ggt Gly													949
40				aaa Lys													997
	_		_	aaa Lys	. –	_			-			_		_	_	_	1045
45				gaa Glu 290													1093
50				acc Thr													1141
				gtg Val													1189
55				gca Ala													1237

5	gaa Glu	gat Asp	gct Ala	gag Glu	aaa Lys 355	gaa Glu	aac Asn	att Ile	gaa Glu	aaa Lys 360	gat Asp	gaa Glu	gat Asp	gat Asp	gta Val 365	gat Asp	1285
	cag Gln	gaa Glu	ctt Leu	gcg Ala 370	aac Asn	ata Ile	gac Asp	cct Pro	acg Thr 375	tgg Trp	ata Ile	gaa Glu	tca Ser	cct Pro 380	aaa Lys	acc Thr	1333
10	aat Asn	ggc Gly	cat His 385	att Ile	gag Glu	aat Asn	ggc Gly	cca Pro 390	ttc Phe	tca Ser	ctg Leu	gag Glu	cag Gln 395	caa Gln	ctg Leu	gac Asp	1381
15	gat Asp	gaa Glu 400	gat Asp	gat Asp	gat Asp	gaa Glu	gaa Glu 405	gac Asp	tgc Cys	cca Pro	aat Asn	cct Pro 410	gag Glu	gaa Glu	tat Tyr	aat Asn	1429
20	ctt Leu 415	gat Asp	gag Glu	cca Pro	aat Asn	gca Ala 420	gaa Glu	agt Ser	gat Asp	tac Tyr	aca Thr 425	tat Tyr	agc Ser	agc Ser	tcc Ser	tat Tyr 430	1477
20															ccc Pro 445		1525
25	tca Ser	cag Gln	ttc Phe	cca Pro 450	gag Glu	ttt Phe	tcc Ser	acc Thr	tcg Ser 455	ttg Leu	ttc Phe	tct Ser	gga Gly	tcc Ser 460	tta Leu	gaa Glu	1573
30			~	_	~ ~		~ ~			~ -	~ -				act Thr		1621
															tca Ser		1669
35															gac Asp		1717
40	-				_	_	_			_	_			_	gta Val 525		1765
															acg Thr		1813
45															gga Gly		1861
50															gca Ala		1909
															gaa Glu		1957
55	tat Tyr	tcc Ser	aga Arg	gac Asp	gaa Glu 595	gac Asp	cac His	ata Ile	gcc Ala	cac His 600	atc Ile	ata Ile	gag Glu	ctg Leu	cta Leu 605	ggc Gly	2005

5	agt att cca agg cac ttt gct cta tct gga aaa tat tct cgg gaa ttc 205: Ser Ile Pro Arg His Phe Ala Leu Ser Gly Lys Tyr Ser Arg Glu Phe 610 615 620	3
	ttc aat cgc aga gga gaa ctg cga cac atc acc aag ctg aag ccc tgg Phe Asn Arg Arg Gly Glu Leu Arg His Ile Thr Lys Leu Lys Pro Trp 625 630 635	1
10	agc ctc ttt gat gta ctt gtg gaa aag tat ggc tgg ccc cat gaa gat Ser Leu Phe Asp Val Leu Val Glu Lys Tyr Gly Trp Pro His Glu Asp 640 645 650	Э
15	gct gca cag ttt aca gat ttc ctg atc ccg atg tta gaa atg gtt cca 219° Ala Ala Gln Phe Thr Asp Phe Leu Ile Pro Met Leu Glu Met Val Pro 655 660 665 670	7
	gaa aaa cga gcc tca gct ggc gaa tgt cgg cat cet tgg ttg aat tet 2249 Glu Lys Arg Ala Ser Ala Gly Glu Cys Arg His Pro Trp Leu Asn Ser 675 680 685	õ
20	tag caaattctac caatattgca ttctgagcta gcaaatgttc ccagtacatt 2298	3
	ggacctaaac ggtgactete attetttaac aggattacaa gtgagetgge tteateetea 2358	3
25	gacctttatt ttgctttgag gtactgttgt ttgacatttt gctttttgtg cactgtgatc 2418	3
	ctggggaagg gtagtctttt gtcttcagct aagtagttta ctgaccattt tcttctggaa 2478	3
	acaataacat gtctctaagc attgtttctt gtgttgtgtg acattcaaat gtcatttttt 2538	}
30	tgaatgaaaa atactttccc ctttgtgttt tggcaggttt tgtaactatt tatgaagaaa 2598	}
	tattttagct gagtactata taatttacaa tcttaagaaa ttatcaagtt ggaaccaaga 2658	3
	aatagcaagg aaatgtacaa ttttatette tggcaaaggg acateattee tgtattatag 2718	}
35	tgtatgtaaa tgcaccctgt aaatgttact ttccattaaa tatgggaggg ggactcaaat 2778	}
	ttcagaaaag ctaccaagtc ttgagtgctt tgtagcctat gttgcatgta gcggacttta 2838	ţ
	actgetecaa ggagttgtge aaaettttea tteeataaca gtetttteae attggatttt 2898	}
40	aaacaaagtg gctctgggtt ataagatgtc attctctata tggcacttta aaggaagaaa 2958	ſ
	agatatgttt eteattetaa aatatgeatt ataatttage agteeeattt gtgattttge 3018	
	atatttttaa aagtactttt aaagaagagc aattteeett taaaaatgtg atggeteagt 3078	
45	accatgtcat gttgcctcct ctgggcgctg taagttaagc tctacataga ttaaattgga 3138	
	gaaacgtgtt aattgtgtgg aatgaaaaaa tacatatatt tttggaaaag catgatcatg 3198	
	cttgtctaga acacaaggta tggtatatac aatttgcagt gcagtgggca gaatacttct 3258	
50	cacageteaa agataacagt gateacatte attecatagg tagetttaeg tgtggetaca 3318	
	acaaatttta ctagcttttt cattgtcttt ccatgaaacg aagttgagaa aatgattttc 3378	
	cctttgcagg ttgcacacag ttttgtttat gcatttcctt aaaattaatt gtagactcca 3438	
55	ggatacaaac catagtaggc aatacaattt agaatgtaat atatagaggt atattagcct 3498	

	ctttagaagt	cagtggattg a	atgtctttt tatt	ttaaat tttaca	ttca ttaaggtgcc 3558
5	tcgtttttga	ctttgtccat t	aacatttat ccat	atgeet ttgeaa	taac tagattgtga 3618
	aaagctaaca	agtgttgtaa c	aataatcca ttgt	ttgagg tgcttg	cagt tgtcttaaaa 3678
	attaaagtgt	tttggttttt t	tttttccag aaaa	iaaaaaa aaaaaa	aaaa aaaaaaaaa 3738
10	ttcctgc				3745
	<210> 2				
	<211> 686 <212> PRT				
15	<213> Homo	sapiens			
	<400> 2 Met Ser Val	Asn Ser Glu 5	Lys Ser Ser S	Ser Ser Glu Ar 10	g Pro Glu Pro 15
20	Gln Gln Lys	Ala Pro Leu 20	Val Pro Pro E 25	Pro Pro Pro Pr	o Pro Pro Pro 30
	Pro Pro Pro		Asp Pro Thr I		o Glu Glu Glu 5
			Glu Glu Gln G 55	lu Asp Pro Al 60	a Asp Tyr Cys
25	Lys Gly Gly	Tyr His Pro	Val Lys Ile (Sly Asp Leu Ph 75	e Asn Gly Arg 80
	Tyr His Val	Ile Arg Lys 85	Leu Gly Trp C	Sly His Phe Se 90	r Thr Val Trp 95
30	Leu Cys Trp	Asp Met Gln	Gly Lys Arg E	he Val Ala Me	t Lys Val Val 110
	Lys Ser Ala	-	Thr Glu Thr A	ala Leu Asp Gl 12	
		_	Ser Asp Pro S		
		Leu Ile Asp 150	Asp Phe Lys I		t Asn Gly Ile 160
<i>35</i>			Glu Val Leu G	Sly His His Le .70	
	Ile Ile Lys		Gln Gly Leu F 185		s Val Lys Ser 190
	Ile Ile Arg		. Gln Gly Leu A 200	sp Tyr Leu Hi 20	
40	Lys Ile Ile 210	His Thr Asp	Ile Lys Pro 0 215	Slu Asn Ile Le 220	u Met Cys Val
	_	Tyr Val Arg 230	Arg Met Ala A		u Trp Gln Lys 240
			Ser Gly Ser A		
45	Gln Lys Pro		Ile Ser Lys A		
,,,	Lys Lys Gln 275	-	Ala Glu Leu I 280	eu Glu Lys Ar 28	g Leu Gln Glu
	Ile Glu Glu 290	Leu Glu Arg	Glu Ala Glu A 295		
50	Ile Thr Ser	Ala Ala Pro 310	Ser Asn Asp G	In Asp Gly Gl 315	u Tyr Cys Pro 320
			Thr Gly Leu G		
	Thr Ala Lys	Asp Asn Gly 340	Glu Ala Glu A 345		
55	Ala Glu Lys 355	Glu Asn Ile	Glu Lys Asp G	lu Asp Asp Va 36	l Asp Gln Glu
	Leu Ala Asn	Ile Asp Pro	Thr Trp Ile G		

		370					375					380				
5	385	Ile				Pro 390					395					400
	_	_			405	Asp				410					415	
				420		Ser			425					430		
10			435			Pro		440					445			
		450				Thr	455					460				
	465	-	-			Leu 470					475					480
15					485	His				490					495	
		_	_	500		Lys			505					510		
20			515			Arg		520					525			
20		530				Cys	535					540				
	545		-		-	Arg 550					555		_		_	560
25					565	Ile	_			570	-				575	
			•	580	•	Leu			585			•		590	•	
		-	595	-		Ile		600					605	-		
30		610				Leu	615	-				620				
	625	_	-			Arg 630				_	635	_		_		640
		-			645	Glu	_	_	_	650				_	655	
35				660		Leu			665					670	GIu	Lys
	Arg	Ala	675	Ala	GIA	Glu	Cys	Arg 680	HIS	Pro	Trp	Leu	Asn 685	ser		
		•														

Claims

40

50

55

- Method for identifying compounds useful for treating and/or preventing Cytomegalovirus infection and/or associated diseases comprising:
 - a) contacting a test compound with one or more cellular kinases selected from the group consisting of RICK, RIP, NIK, MKK3, and SRPK-2; and
 - b) detecting a change in activity of said cellular kinase.
 - 2. Method for detecting Cytomegalovirus infection and/or associated diseases in an individual comprising:
 - a) providing a sample from said individual; and
 - b) detecting activity, in said sample, of one or more cellular kinases selected from the group consisting of RICK, RIP, NIK, MKK3, and SRPK-2.
 - 3. Method for detecting Cytomegalovirus infection and/or associated diseases in cells and/or cell lysates comprising:

a) providing a sample from said cells; and

10

25

40

45

- b) detecting activity, in said sample, of one or more cellular kinases selected from the group consisting of RICK, RIP, NIK, MKK3, and SRPK-2.
- A monoclonal or polyclonal antibody that binds to a cellular kinase selected from the group consisting of RICK, RIP, NIK, MKK3, and SRPK-2.
 - 5. Method for preventing and/or treating Cytomegalovirus infection and/or associated diseases in an individual by administering a pharmaceutically effective amount of an inhibitor to said individual, wherein said inhibitor inhibits at least partially the activity of one or more cellular kinases selected from the group consisting of RICK, RIP, NIK, MKK3, and SRPK-2, or wherein said inhibitor inhibits at least partially the production of one or more cellular kinases selected from the group consisting of RICK, RIP, NIK, MKK3, and SRPK-2.
- 6. Method for regulating the production of Cytomegalovirus in an individual by administering an individual a pharmaceutically effective amount of an inhibitor wherein said inhibitor inhibits at least partially the activity of one or more cellular kinases selected from the group consisting of RICK, RIP, NIK, MKK3, and SRPK-2, or wherein said inhibitor at least partially inhibits the production of one or more cellular kinases selected from the group consisting of RICK, RIP, NIK, MKK3, and SRPK-2.
- 7. Method for regulating the production of Cytomegalovirus in cells by administering the cells a pharmaceutically effective amount of an inhibitor wherein said inhibitor inhibits at least partially the activity of one or more cellular kinases selected from the group consisting of RICK, RIP, NIK, MKK3, and SRPK-2, or wherein said inhibitor at least partially inhibits the production of one or more cellular kinases selected from the group consisting of RICK, RIP, NIK, MKK3, and SRPK-2 in the cells.
 - 8. Method according to claim 5, 6, or 7 wherein the inhibitor is a monoclonal or polyclonal antibody which binds to a cellular kinase selected from the group consisting of RICK, RIP, NIK, MKK3, and SRPK-2.
- 9. Method for preventing and/or treating Cytomegalovirus infection and/or associated diseases in an individual by administering a pharmaceutically effective amount of an activator to said individual, wherein said activator activates at least partially the activity of one or more cellular kinases selected from the group consisting of RICK, RIP, NIK, MKK3, and SRPK-2, or wherein said activator activates or stimulates at least partially the production of one or more cellular kinases selected from the group consisting of RICK, RIP, NIK, MKK3, and SRPK-2.
- 10. Method for regulating the production of Cytomegalovirus in an individual by administering an individual a pharmaceutically effective amount of an activator wherein said activator activates at least partially the activity of one or more cellular kinases selected from the group consisting of RICK, RIP, NIK, MKK3, and SRPK-2, or wherein said activator at least partially activates or stimulates the production of one or more cellular kinases selected from the group consisting of RICK, RIP, NIK, MKK3, and SRPK-2.
 - 11. Method for regulating the production of Cytomegalovirus in cells by administering the cells a pharmaceutically effective amount of an activator wherein said activator activates at least partially the activity of one or more cellular kinases selected from the group consisting of RICK, RIP, NIK, MKK3, and SRPK-2, or wherein said activator at least partially activates or stimulates the production of one or more cellular kinases selected from the group consisting of RICK, RIP, NIK, MKK3, and SRPK-2 in the cells.
 - 12. Oligonucleotide that binds to the DNA or RNA encoding a cellular kinase selected from the group consisting of RICK, RIP, NIK, MKK3, and SRPK-2.
- 13. Method for regulating the expression of at least one cellular kinase selected from the group consisting of RICK, RIP, NIK, MKK3, and SRPK-2 in an individual comprising the step of administering the individual a pharmaceutically effective amount of an inhibitor wherein said inhibitor inhibits at least partially the transcription of DNA or the translation of RNA encoding one of said cellular kinases.
- 14. Method for regulating the expression of at least one cellular kinase selected from the group consisting of RICK, RIP, NIK, MKK3, and SRPK-2 in cells comprising the step of administering the cells a pharmaceutically effective amount of an inhibitor wherein said inhibitor inhibits at least partially the transcription of DNA or the translation of RNA encoding one of said cellular kinases.

- **15.** Method according to claim 5, 6, 7, 13, or 14 wherein the inhibitor is a oligonucleotide which binds to the DNA and/ or RNA encoding a cellular kinase selected from the group consisting of RICK, RIP, NIK, MKK3, and SRPK-2.
- 16. Method for regulating the expression of at least one cellular kinase selected from the group consisting of RICK, RIP, NIK, MKK3, and SRPK-2 in an individual comprising the step of administering the individual a pharmaceutically effective amount of an activator wherein said activator activates at least partially the transcription of DNA or the translation of RNA encoding one of said cellular kinases.

5

20

25

45

50

- 17. Method for regulating the expression of at least one cellular kinase selected from the group consisting of RICK, RIP, NIK, MKK3, and SRPK-2 in cells comprising the step of administering the cells a pharmaceutically effective amount of an activator wherein said activator activates at least partially the transcription of DNA or the translation of RNA encoding one of said cellular kinases.
- 18. A solid support useful for detecting Cytomegalovirus infection of an individual comprising one or more immobilized oligonucleotides, wherein said oligonucleotide(s) is (are) capable of detecting activity of one or more cellular kinases selected from the group consisting of: RICK, RIP, NIK, MKK3, and SRPK-2.
 - 19. A solid support useful for detecting Cytomegalovirus infection of cells comprising one or more immobilized oligonucleotides, wherein said oligonucleotide(s) is (are) capable of detecting activity of one or more cellular kinases selected from the group consisting of RICK, RIP, NIK, MKK3, and SRPK-2.
 - 20. A solid support useful for screening compounds useful for treating and/or preventing Cytomegalovirus infection comprising one or more immobilized oligonucleotides, wherein said oligonucleotide(s) encode one or more cellular kinases selected from the group consisting of RICK, RIP, NIK, MKK3, and SRPK-2.
 - 21. A solid support useful for screening compounds useful for treating and/or preventing Cytomegalovirus infection comprising one or more immobilized cellular kinases selected from the group consisting of RICK, RIP, NIK, MKK3, and SRPK-2.
- 22. Composition useful to treat an individual afflicted with Cytomegalovirus and/or associated diseases comprising one or more inhibitors capable of inhibiting activity of one or more cellular kinases selected from the group consisting of RICK, RIP, NIK, MKK3, and SRPK-2 or capable of decreasing the expression of at least one cellular kinase selected from the group consisting of RICK, RIP, NIK, MKK3, and SRPK-2.
- 23. Composition useful to treat an individual afflicted with Cytomegalovirus and/or associated diseases comprising one or more activators capable of increasing activity of one or more cellular kinases selected from the group consisting of RICK, RIP, NIK, MKK3, and SRPK-2 or capable of increasing the expression of at least one cellular kinase selected from the group consisting of RICK, RIP, NIK, MKK3, and SRPK-2.
- 40 24. Composition useful to treat an individual afflicted with Cytomegalovirus comprising at least one compound selected from the group consisting of
 - 6-(2,6-Dichlorophenyl)-8-methyl-2-(3-morpholin-4-yl-propylamino)-8H-pyrido[2,3-d]pyrimidin-7-one;
 - 8-methyl-6-phenyl-2-(pyridin-4-yl-amino)-8H-pyrido[2,3-d]pyrimidin-7-one;
 - 6-(2,6-Dichlorophenyl)-8-methyl-2-[3-(4-methylpiperazin-1-yl)-propylamino]-8H-pyrido[2,3-d]pyrimidin-7-one;
 - (3-Bromophenyl)-(6,7-dimethoxyquinazolin-4-yl)-amine;
 - (3-Bromophenyl)-(6,7-diethoxyquinazolin-4-yl)-amine;
 - 2-(3,4-Dihydroxyphenyl)-3,5,7-trihydroxychromen-4-one;
 - 5-Cloro-3-(1H-pyrrol-2-ylmethylene)-1,3-dihydroindol-2-one;
 - 4-Quinolin-4-ylmethylene-4H-isoquinoline-1,3-dione

and/or phamaceutically acceptable salts of these compounds.

- 25. Composition according to any one of claims 17 19 further comprising pharmaceutically acceptable carriers, excipient, and/or diluents.
 - 26. Use of the compounds selected from the group comprising:

6-(2,6-Dichlorophenyl)-8-methyl-2-(3-morpholin-4-yl-propylamino)-8H-pyrido[2,3-d]pyrimidin-7-one; 8-methyl-6-phenyl-2-(pyridin-4-yl-amino)-8H-pyrido[2,3-d]pyrimidin-7-one; 6-(2,6-Dichlorophenyl)-8-methyl-2-[3-(4-methylpiperazin-1-yl)-propylamino]-8H-pyrido[2,3-d]pyrimidin-7-one; 5 (3-Bromophenyl)-(6,7-dimethoxyquinazolin-4-yl)-amine; (3-Bromophenyl)-(6,7-diethoxyquinazolin-4-yl)-amine and pharmaceutically acceptable salts of these compounds as an inhibitor of the cellular kinase RICK. 10 27. Use of the compounds selected from the group comprising: 2-(3,4-Dihydroxyphenyl)-3,5,7-trihydroxychromen-4-one; 5-Cloro-3-(1 H-pyrrol-2-ylmethylene)-1,3-dihydroindol-2-one; 4-Quinolin-4-ylmethylene-4H-isoquinoline-1,3-dione; and 15 pharmaceutically acceptable salts of these compounds as an inhibitor of the cellular kinase RIP. 28. Use of a compound according to claim 26 or 27 for the manufacture of a pharmaceutical composition for prophylaxis and/or treatment of Cytomegalovirus infection and/or diseases associated therewith. 20 25 30 35 40 45 50 55

Figures

Fig. 1

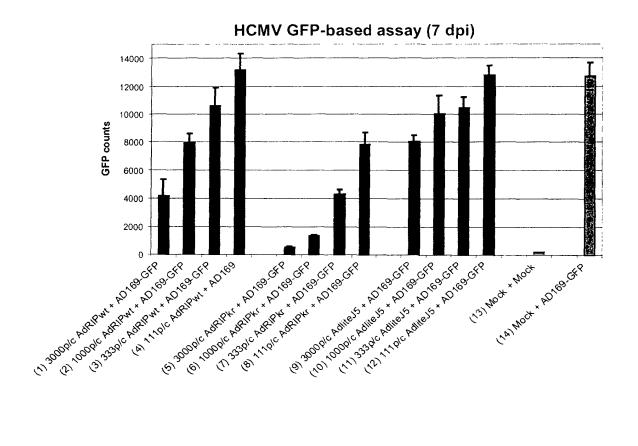


Fig. 2

